

WO 00/28040

PCT/US99/26260

## SEQUENCE LISTING

&lt;110&gt; Wong et al.

&lt;120&gt; CLONING OF RHADINOVIRUS GENOME AND METHODS FOR ITS USE

&lt;130&gt; 53683

&lt;140&gt; 60/107,507

&lt;141&gt; 1998-11-06

&lt;140&gt; 60/109,409

&lt;141&gt; 1998-11-20

&lt;160&gt; 179

&lt;170&gt;

&lt;210&gt; 1

&lt;211&gt; 133719

&lt;212&gt; DNA

&lt;213&gt; Macaca mulatta rhadinovirus 17577

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13021	TTCGTGGGCG	ACGCCATCTC	CGTGACGGAC	TGCGTGGCGG	TGGACCAGGC	GTCCGTACAG
13081	ATCCACAAGA	GCCTCCGCAC	GTCCACCCCG	GGGATCTGCT	ACTCGCGCCC	CCCGGTCACG
13141	TTCAGGTTCC	TCAACAGCAC	CACGCTGTTC	AAGGGCCAGC	TGGGACCCAG	AAACGAGATC
13201	ATACTGACGG	ACAACCAGGT	GGAGGCGTGC	AAAGAGACGT	GCGAACACTA	CTTCATAGCG
13261	AGCAACGTAA	CCTACTACTA	CAAAGACTAC	GTCTTCGTGA	AAAAAATTAA	CACCTCCGAG
13321	ATATCCACCC	TCGGTACGTT	CATCGCCCTG	AACCTGTCGT	TTATAGAGAA	CATAGATTTT
13381	AGGGTCATCG	AGCTGTACAG	CCGCGCGGAG	AAAAAGCTGT	CCGGGAGCGT	TTTCGATATA
13441	GAAACCATGT	TCAGGGAATA	CAACTACTAC	ACGCAACGCC	TGGCGGGACT	CCGGGAGGAC
13501	CTGGACAACA	CGATCGACCT	GAACCGCGAC	CGCCTGGCCC	GCGACCTGTC	CGAGATAGTC
13561	GCGGACCTGG	GCGATGTCGG	CCGCACGGTC	GTTAACGTGG	CCAGTAGCGT	GATAACCCTG
13621	TTCGGATCAA	TCGTGAGCGG	GTTCAATTAAC	TTTATAAAGA	GTCCGTTCCG	GGGCATGCTC
13681	ATGATCCTGG	TGATTGTGGC	GGTCGTCTTG	ATCGTGTTCG	CGCTAAACCG	GCGACCAAC
13741	GCCATCGCCC	AGGCCCCCAT	CAGGATGATC	TACCCCGACA	TAGACAAAT	CGCGCCCTCT
13801	GGCGGTAAAG	TCGACCAGGA	CGAGATTAAA	AACATTCTCG	CCGGCATGCA	CCAGCTACAG
13861	CAGGAAGAGC	GTAGGCGGTT	AGAGGAAACG	CAGAGGTCAG	CGCCCTCGCT	TTTCCGGCGC
13921	GCGTCAAGCG	GACTAAAACG	TCGCTTTAGG	GGATATAAAC	CGCTGGAAAA	CGAAGAGGCT
13981	CAAGAGTATG	AAATGAGCAA	ATAACCACAC	CCACACGCCT	GTACTTGCCG	CCCGCCAGAG
14041	CCGCGCGGCC	AATCGATTTC	CGCACCCGGC	CGGTCCCCGA	CACCCTCTAG	CGCCCCCGGG
14101	CGTCCGCGGT	CTGTATCAAT	CATGGATTTT	TTTAACCCGT	ACCTGGGCCC	TCGCGGACCA
14161	CGCCCCCACT	CACACAGAGG	CACCGATGCT	CCCGCCCCCT	CCGGCGCCGG	AGCCGGTTCAG
14221	CCGCCACCAG	ACGTTTGCAG	GCTCATCCCC	GCCTGCCTCC	GAACGCCAGG	GGCAGGCGGG
14281	ATGATCCCCG	TCACGATCCC	GTTCCCGCCA	ACGTACTTCG	AGAACGGTGC	TCGCGGAGAC
14341	GTGCTGCTCG	CCAACGAACG	GTCCATGTGG	ACGGCGCGCG	ACCGCAAGCC	CGTCGCCCCG
14401	GACCCCAAG	ACCAATCCAT	CACGTTTCAC	GCGTACGACG	TCGTTGAAAC	AACGTACGCG
14461	GCGGACAGGT	GTGCCGAGGT	ACCTAGCCGC	TTCCAAACGG	ACATTATCCC	AAGCGGAACC
14521	GTGCTCAAGC	TCCTGGGGCG	AACCGAGGAC	GGCACCAGCG	TGTGCGTGAA	CGTGTTCCGT
14581	CAACAGGTAT	ATTTCTACGC	GAAGGTTCCA	GCCGGCATT	ACGTCACCCA	CATCCTCCAG
14641	CAGGCCCTCA	AGAACACAGC	CGGCCGTGCC	GCGTGCGGCT	TCTCGACCAG	AAGAGTGAAC
14701	AAAAGAATTC	TCAAAACGTA	CGACGTCCCG	GAGCATCCCC	TCACGGAAAT	CACGCTATCG
14761	TCCGGTTCCA	TGCTCTCGAC	CCTCAGCGAC	CGCCTCGTCG	CGTGCGGGTG	CGAGGTGTTT
14821	GAGTCAAACG	TGGACGCCGT	TCGCCGGTTC	GTTCTGGATC	ACGGGTTTAC	CACGTTCCGG
14881	TGGTACTCGT	GCGCGCGCGC	CACGCCCCGC	CTGGCGGCCA	GAGATGCCAG	GACGGCCCTG
14941	GAGTTTGACT	GCAGCTGGGA	GGACCTCAGC	GTTCAAGCGG	ACCGCAGCGA	CTGGCCCCCG
15001	TACCGCATCG	TGGCCTTTGA	TATCGAGTGC	ACTGGAGAGG	CGGGATTTCC	GTGCGCCACG
15061	CGCGACGGCG	ACGCGGTGAT	CCAGATCTCC	TGCGTCTTCT	ACACGACCAG	GGAAGGCGCG
15121	CCCAATCCGC	CAAACATACT	GTTCAGCGTC	GGGACGTGCG	ACCCCATCCC	GGACACCGAC
15181	GTTTTGGAGT	TTCCGTCGGA	ATATGACATG	CTGGTGTCGT	TCTTCGCCAT	GATCCGCGAC
15241	TTCGAGGTGG	ACTTTTTTAAC	CGGCTATAAC	ATCTCAAAC	TCGATCTCCC	GTACCTAATC
15301	ACGCGAGCGT	CCCAGGTGTA	CAACCTTCGA	TTAAACGAAT	ACACAAAAAT	AAAAACCGGC
15361	TCCATCTTTG	AAGTTCACGA	GCCCCGTGGC	GGGGGAGGGG	GTTTCATGAG	GTCGGTCTCA
15421	AAAATTAATA	TAGCGGGCAT	CGTCCCCATA	GACATGTACC	AGGTGTGTCG	CGAAAAGCTC
15481	AGCCTCTCCG	ACTACAAACT	GGACACGGTG	GCCAGGCAGT	GTCTGGGTGG	GAAAAAGAG
15541	GACGTATCGT	ACAAGGACAT	TCCCCCTCTG	TTTCGCTCAG	GTCCGGGCGG	CAGGGCTAAG
15601	GTGGGACGCT	ATTGCGTGAT	GGACTCGGTC	CTGGTGATGG	ACCTCTTAAA	AATGTTTATG
15661	ATACACGTGG	AGATTTTCGGA	GATAGCCAAG	CTGGCCAAGA	TTCAGGCCAG	GCGCGTCTCT

15721	ACGGACGGCC	AACAGCTCCG	CGTGTCTCTCC	TGCCTGCTGG	AGGCCGCGGC	CAGGGAGAAC
15781	TTTATCCTCC	CGGTTCCAAC	GCCCCAGGGA	CAGGGGGGCT	ATCAGGGCGC	GACGGTGATC
15841	AACCCCATTC	CGGGGTTTTA	CGACGAGCCG	GTCCTGGTGG	TCGATTTTGC	CAGCCTGTAC
15901	CCGAGCATCA	TCCAGGCGCA	CAACCTGTGC	TACTCCACCA	TGATACACGG	ACGAGACCTG
15961	CACCTGCACC	CCAACCTGAC	GCCGGACGAG	TACGAGACGT	TCGTGCTGAG	CGGCGGACCG
16021	GTACATTTTG	TAAAAAACA	CAAGCGGGAG	TCTCTGCTGG	GAAGACTGCT	AACCGTGTGG
16081	TTAGAAAAGC	GAAGGGCGAT	CCGGCGCACC	CTGGCGGCGT	GCGATGACCC	GTCGCTAAAA
16141	ACCATCTTAG	ATAAACAACA	GCTGGCCATC	AAGGTGACAT	GTAACGCGGT	TTACGGGTTC
16201	ACCGGGGTGG	CCAGCGGCCT	CCTCCCATGC	ATTAACATAG	CGGAAACCGT	GACGCTCCGG
16261	GGGCGCACGA	TGCTGGAGAT	GTCAAAGTCT	TACGTGGAGG	CCCTGACGAC	GGAAGACCTG
16321	CGAACGCGTC	TCGGTCGCGA	GGTGACCGCC	CGTCACGGCG	CGCGGTTTCG	CGTCGTCTAC
16381	GGTGACACCG	ACTCCCTCTT	TATCGCGTGC	GACGGTTATT	CCGCGGAAGC	CGTTTCCGCT
16441	TTCTGTGACG	ATCTGGCCGC	CAGGATCACT	GCGGACCTGT	TCCCCCACC	CATTAAGCTA
16501	GAGGCGGAAA	AGACGTTCAA	GTGTCTGCTG	CTGCTGACGA	AAAAGCGCTA	CATCGGGGTC
16561	CTATTGAACG	ACAAAATGGT	CATGAAAGGG	GTCGACCTCA	TTCGCAAAAC	GGCCTGCAAG
16621	TTTGTCCAGG	AGCGATGCCG	CGCCATCCTG	GACCTGGTGC	TCCACGATCC	GGAGGTCAAG
16681	GCTGCGGCGC	GGCTGTTGTG	CAAGCGGCCG	CCGCACGCGG	TATACGAGGA	GGGGCTGCCG
16741	GCTGGCTTTA	TAAAAATCGT	AGAGGTCTCT	AACGCGAGCT	ATCTGGACCT	CCGAAACAGC
16801	GTCGTGCCCA	TCGAGCAGTT	AACGTTCTCC	ACCGAGCTCA	GCCGCCCGGT	CTGCGATTAC
16861	AAGACCACCA	ACCTGCCCCA	CCTGGCGGTG	TACCAAAAGC	TGGCGAGCAG	GTGCGAGGAG
16921	CTGCCCCAGG	TGCACGATAG	AATCCCCTAC	GTGTTCTGTT	ACGCGCCCGG	GTCCCTAAAG
16981	TCGGACCTGG	CCGAACACCC	GGATTACGTC	AGACAGCACC	AGATTCCTCG	CGCGGTCGAC
17041	CTATATTTTC	ACAAACTGGT	GCACGGCGCG	GCCAACATCC	TCCAGTGTCT	GTTCCGGCAAC
17101	AACGCGGACA	CCACGGTGGC	CATCCTCTAC	AATTTTCTCA	ACGTCCCGTA	TAAGCTGTTC
17161	TCGTGAACGC	CAATTGGAGA	ACGCCAACAT	AAGACGCCGC	GCCAGCGGAG	TCCGAGGGGA
17221	GAGCTCGAGC	GGCGAGGAGC	GACCAACGGA	GACCGCCACC	ATGCTGGTTA	ACGAACTGTC
17281	GGTGGTCTCT	GGCGACTGGG	AGGTGACTTT	TCACCGGGGT	AGATTTCAGCT	TCGTCAACCT
17341	CACCCGCTTG	CAAACGTTCA	AGGGCCACGG	GGGCTACGCC	AGGGTCCGAC	TCCCCTTCTC
17401	GCTCGACCAG	TTACTCCACC	AACATTTTCG	GTTTCGGACTC	GTGACGCGTC	TCAAGGAACT
17461	GCCCCCTTTC	TCCGACTGCG	TGGCCCTTAT	CGCCCCGTTG	GATTCCGGCG	GCGACGCGGA
17521	CGCGGCGCGC	GTGGCCCCCG	GGTTCGTGCT	GGACTCCTCT	CGCCCGCTGA	CGGTGTGGGT
17581	AAACGCGAGC	GGGCGGCACA	CGATCCGGTT	CTGCCTCCTC	TTTCTAAAGC	CGATCGGACCT
17641	GGAGCGCGCG	GTCACGTACG	TCTTCGGCGG	GAACGGCGGC	GCGCGCTCGG	AGGGCACCCC
17701	AAAGCCCACC	TGCGCGACCG	AAAGCCTGCC	CGGTGGACCC	CTGCGCGTCT	CCGGCGAGGC
17761	GTCTCAGACG	TCGCCCCATT	CTTTCGTTGC	GTATTTTCCC	ACGGCCAACT	CGGTGGCCTG
17821	CCTAAGCCTG	TTGCGGTTAC	AGGTGAGGCC	GTTTTCGGAT	GACGCGGCGC	ACAGGGACGC
17881	GCGGATCTCC	CCGAAATACG	TCACGTTTAG	TAACCTCCGG	GGTAACGTCT	GCAAGGCGTC
17941	CGTTCACACG	CTGTCCCCGT	CGCGGTGTAA	AACGGCGCAA	ATGGAAATCA	TCTACGCTCC
18001	CGGGGACCCC	AACGCCGAGA	TAGTCCTGGG	CCAGTCCGGA	CCCGTCCTGC	CCACCCACAC
18061	CGGCGGCCGC	GTATTGGGGG	TCTACGCCGA	CGCCGAAAAA	ACCATCCAAC	CTGGAAGCTC
18121	CGCGGAAGTC	CGGGTTTCA	TAATCTTCCA	ACAGGGAGCG	GCCGCTCGGG	GCGATCTGGC
18181	GTTTCTGGTC	ACGGGCGTGG	CACCGGAGCC	CCTATTCTGC	GTCACCCCGG	CACCTCTTGT
18241	TTCCGGTTTG	ACAACCCACC	TGCGCCTATT	CAACCCCAAC	GGTACCCCCA	CGACTATAAA
18301	AAGAGACACC	CTTGTGGCCG	CCGCCGCGCC	CTGCCCCGTG	GTGCGATTAA	GCTCCGCCGA
18361	CGACGCGCCG	CGAGACCTCG	TCGCGTCACC	AGACACCGGG	GCGCTCTCCA	TTAACGCGTT
18421	CACAATCCCG	GTCGGTTTCC	CAGGGGTGGT	CTCGGCGGAG	TGTCACGTGT	CCCTACGCGA
18481	CAACGGGGTC	CACGAACGCA	TGAACCATTG	ACGGCAACGA	TGGGAACACC	GGTGCGTTTC
18541	TTTCGCGGCG	AGTGGCAGAC	CTCGAGTCTA	GTGGACAACG	GCACGCCACG	GTACAGCTCC
18601	CTGGTGTGGG	CCGCCACTAT	TCACGACGGC	TACCTGACAC	TGGTGAACAG	GTCAGAGCTG
18661	TGCGTCACGG	AGAGGTCTCC	GTGTCTGCCG	GCATGCCCCA	GCATCGGGAG	ACTGGTCGGG
18721	AAGAGGTTTC	CCGGCTTCGC	CTTTGCCAGC	GCCACTCTGG	GCGATCGGGG	AACACGCACC
18781	GTGTTCTACG	CGTTCGGTCA	CCTGCGGATT	CCACTGGACA	TAGTACCCGC	CGTGGTCGAG
18841	CGCGCGGATC	GCGAGCTGGT	CTGCGGGT	CACGCTCCGC	AAACAACGCG	GGTGTGCGGA
18901	TACGGACTTA	AGGTATTCGT	GGCGATCGTT	ACGGTGGTGC	GCCCGCCCGG	GGTGTTCCTA
18961	CACTTTCCAC	AAGACCGCGT	TCCGATCGCG	CTGACAGACG	CGTGCAGCCA	GGAGGGCTCC
19021	AGGCTAACCT	CTGAAGAGCC	GTGGATAAAA	ATTCAAGGCT	TTCCCGTCTT	ATCTGACGAG
19081	ACCGCGCACC	CATTTCTCTT	AACCCAGAAG	ACCAAGCCCT	TTACCGAGCG	AAAGTTTTGC

19141	CGCCTGATCA	TGGACAACGA	CCAGCGCAGC	GCCGTCAACA	CCGTCTACCT	GGGAAAGCAG
19201	CACGTGAGGG	TGACCGTGAC	CCGCCCCCGG	GAAACAATCG	TCACCGACGG	CCCCGTGACG
19261	GCGACCCGTG	CCCTCACCGG	TAATGCGCCA	ATCGCCTTTC	GCCACAACCC	ATACTTTGAA
19321	CTCCCGTGTT	CGTCCACAAC	GGCGATATTC	ACGCCCGTGG	TGTACGTGGG	CCTGACCGTG
19381	TGCATCCAC	CCAAGTGTAG	CAAATTCGTA	AGGTACGGTA	ACACCTACGT	CTCGGCATTT
19441	AACCGCAAGC	TGACGGCGAT	TATTAGCAAT	CACGCCACA	ACGGCGGGTT	CCGGATTGAG
19501	GAATGCGAGT	GGCCACCGAA	CCGGGAGATA	GAGATTTTGG	TAACCAACGT	GTCCAGGCC
19561	CCGGTGTACA	TCAGCACCGG	GACGCAGCTG	GGGCAAGCCA	TCTTCGTGTT	CGCGCCGCGG
19621	TTCGGTGGCC	CGGCGAAACT	GCGGCAGCTC	CTCGGCCACC	GATCGCGCGC	CCTGGAGCTG
19681	CCGGGCGGGG	TGACAGTGGA	CAGCCAAAAA	CTGTGTAGGT	TTGAGACCAT	GTACCTGTTT
19741	TCCACGTAAA	TTACTAATAA	ACCGTTTGCT	CGTATCGCTC	ACACAACGCC	AAACCGTCTC
19801	TCATTCTCGG	GGTCGCGCGC	CTCGCGAACA	CACAAGGTGG	CTCAAACACC	CCCCTCCGCG
19861	ACCTCGCCA	CACAAAACCA	GTTAACGCCT	TCCGTTAGAT	GCAGTTTATT	TATTATTTTA
19921	TTACATCATA	GCTATTGCGC	GGCGCCCGTC	CCGCAAAAAC	ATCTGTAGAT	ATTCCAGTAT
19981	GCGAAACGCG	CTGAGAACAA	CGTCCCGGGG	GCTCTGCCCT	CCCAACGCAC	GCACGGTTTT
20041	TTCATTAGAC	TCCGCGCCAC	CTATCTTGTT	ATTTACGGGA	AGCTCCTCAA	TTAGAGAGTC
20101	GAGGGCGGAC	AGCACCACGG	TCACCGCGGA	GCCAATGGCC	GCGGCGTCCG	CGTCCCCGGG
20161	CGCGTCGTCC	AGCAGCATCC	TCAGTCGCGT	CAGGTAAGTC	TCGTAGGCCG	GGAGCCCGCG
20221	AACCATGGCG	TTCATACACT	CGGCCCCGCG	AAACTCGCGG	CGCTGGCACT	CAACGTTAGA
20281	CATCAATATA	GGGAAGTGAA	CAAACGACAT	GATGCGCGGG	TACTCACGCA	CCCGACAGAG
20341	AGTGGAGTGG	TGGCACAGGT	AAACGACCA	GCGCTGTATG	TTTAACCCCA	CTTCAGACAG
20401	GCGCCCCCTG	GTCCTGTCAA	GAGAGGCGCG	GTTGCCGGCC	CACTGGGGGA	GAACGTTAAT
20461	TCCAGCGGAG	GTCGGGGGAG	GCGCCAGCGT	AGGGCTGGCC	GCCCAACACG	ACAGGTAAAA
20521	CAAGACGAAC	CAGACAGGGA	ACATGACCGG	TTAAATTACT	CGCTTACAAT	CGCGGGCGGC
20581	GGCCGGTCAA	CGCCAGGTCC	ATTAATAACA	CACCGGCCGC	CAACCCCCAA	CGCGGGGCCG
20641	CGCCCTGGAA	CGCGGTTCCT	TCCAATCGCA	AAGAACC CGG	TCACAAAAG	GGCTCGTTTT
20701	GAACCCATTT	TGTGCCATCG	GGTTTCGTTT	TCAGATACGG	AAACGGCTCG	TCCAAAACA
20761	CCCAACGGGG	GTGTGCTCAA	ACGGCCATCT	CCATCTCTAT	GTGGGGATGG	GGTCTGTAGC
20821	CCTCGAGACT	CAGATCCGCG	CGCGTAAAGT	CCTCCAGACG	CGCCACTTT	CTCAAATCT
20881	TCAGCCGCGG	AAACGGACGC	GGGGTCCTCC	GCAGCTGAAG	CAGCAGGGGA	TCAACGTGGT
20941	TGTTGTAAAC	GTGGGCGTCA	CCCAAGGTGT	GCACAAAGTC	TCCCGGGGTC	AGGCCCGTGA
21001	CGTGAGCGAT	CAGATACGTC	AGGAGGGCGT	AGCTGGCGAT	GTAAACGGG	ACCCGAGGC
21061	CCATGTGCGC	GGACCTCTGG	TACAGCTGGC	AGGACAGCTC	CCCCCGAGCC	ACGTAAACT
21121	GACACAAAAC	GTGACAAGGA	GGGAGAGCCA	TCCGCGCGAG	GTCCGCGGGG	TTCCACGCGC
21181	ACATAACGAT	GCGCCGATCG	TGGGGCCGCC	TGTTAATTAG	ATCCACCACG	TAACGCAGCT
21241	GGTCCACCCC	CTGACCCTCG	TAGTTGGCGT	CGGCCCCCCT	GTACTCCGCC	CCAAAATGTC
21301	TCCACTGGAA	CCCGTACACC	GGCCCCAGAT	CGCCCTCGCG	GCGGTCCCGG	AAGCCCTGCG
21361	CCGCCAAAAA	GGCGCGGGAC	CCGTGCGCGT	CCCAAATTTT	TACGCCGCGG	CGCGACAGTT
21421	CGGTGGAGTC	GGTGGAGCCC	CTGATAAACC	ACAGCAACTC	CTCCACGACG	CCCCTCCAAA
21481	ACACCCTTTT	GGTGGTTAAC	AGAGGAAACT	CGTCCCTGAG	GTTATATCGG	GCCTGAAGCC
21541	CGAACACGGA	CCTGGTGCCC	ACGCCCGTCC	TGTCCTCCCT	CTGCACGCCG	TGTTTAATAA
21601	TTAAATCCAG	GTGCGCCAAG	TACTGCAGCT	CGCCGTGGTC	GCCGCGGCAC	GCGCAGCGGA
21661	CGGCCGCGGC	ACCTTCGGGT	GCGGAAAAAA	CCCGGAGTCG	TGCGGCCCGG	ATTCCAGCCA
21721	CGCAAACGGG	TATAATTTTT	TTAACGTAGC	AGATAACGAG	ATGCACCAGG	ACAATCATAA
21781	TTACAACCGA	GGTGCTTTAC	AACTATAACC	GTAACGGGCT	GAAGACGAAA	CTTATTTTAA
21841	AGGCAGCTTG	GGCGGGGCGG	ACCCAGGCGG	CGGCCCGGCT	GTTAGCGAGT	AATAAACCCA
21901	CGTGGTCTCC	CCAAGCGTCT	CCCCGAGCGT	GAGCGTCTCC	CCAAACGCCA	AAAACAACAC
21961	AGATAATAAA	ATAAATAACA	TGTTTATTTT	TTATAAACTT	AACGCGCGAT	TATGCTTCTT
22021	AACAGGGGCC	AATAGAAGCA	ACGTGCGAGG	CGGCATTGCG	GCGCTCAAAG	GCAACCGCAA
22081	CACCGGCGCT	CGTAACAGCG	TAAACAAACA	ACTATTAGCC	ATTCGTAACC	GTAAATCCA
22141	ACCCTCTGCG	TCCCGAGCCA	TTCGCCCAGC	TGCGTCTACG	TCGAGGGTGT	TTCTAAGTCG
22201	CATTTTGAGC	CCTTGGCAGC	GCCGCGTCTG	CGTTAATGGT	GTTTCTAAAT	CCCGCTGCCA
22261	AGGCCCTCTT	CGGAATCGTC	AATCAGGCTG	CGCCGGGTTT	TTCTGCCCTT	TTTGGGACGC
22321	TTGTCCACCG	CCTGAAGCAG	TTTCTTCACT	GCGTCGTCCC	CGGGATTGGC	ACACAGCTTT
22381	TTACCGCGGC	GAGTGTTTAA	TATCACGGCG	TCCACCGAGC	ACTGCGACGA	GGTGTGGGAG
22441	TAAGAGACCA	CTAAACGGGG	TGGCGGCAGA	TGAGTTACAT	ACCCCAAACA	CGAGGTTTCG
22501	GGCGCGGGTC	CGCTCATGGA	GCCCATAGGA	AAGGCATAAT	CAACTACACA	CGCGAACACC

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22561	GCAAAAAAAAA	CGCACACGAA	AAGGCCCTC	ATGGTGACAA	AAACACAAAA	CAAAGACGGG
22621	GTCGCTCGTG	CGTTGCCACC	AGCGCCACGC	TCGCCGTGCT	CGGCCACGGT	TGACGGCGTC
22681	CATTAATACC	CGCGGCCCGG	GCCCGCCTCG	ACGGAACCA	GGCGTGGTTA	GCCACCTGAC
22741	GCACGTGCCT	CCCGTCCAGA	TGTGGCATTG	CGAGTCCCAA	AACGGGCGTG	TCTAACTACG
22801	GTCGCCCCGG	GGTTCAAAGA	ACGGGGATCC	CCCAAACAGG	TAAAAAGCTT	TTTGTCCAAA
22861	CAGAAAGCCA	ACAGCAACCG	AAAGGAATCT	CCAAAACAAG	CAAGGGTCCC	GTGCCGGCGA
22921	TAAACGAAA	ATTTCCCTCA	CCTAAACACA	CGCGCCGCGT	AAACTGTAA	AACACGCAAT
22981	ACTTCCTAGG	CCTGTGTTTA	ATAAAACACA	CAAGGGTGT	AATTTGCGCG	GGCCCTTTGG
23041	GCCCCGCGCG	TCTGGGGGTG	CGTTGTCTCG	TTGTGGTTCG	CAATCACCAC	CCCGCCTGCC
23101	CGAAAACAGG	GCAACAACCC	CCTGGCCTAG	TTTTTTAAAA	ACTTAACACC	GGCAAGGGGA
23161	GAGGGAGAAG	GGGTGCGGCT	AAATGGGCTG	TTAGCAGCCA	ACCGCAGCGG	GGAAAAAAGG
23221	GGGCAACGCG	CGGGTTAACA	CAAACAGCAA	CAGGCGCCAA	ACCCAATATA	ACACAACGCA
23281	TATTCCTGTG	GTTTCGCGGG	CCGGGGTCGG	TACGGCTAAC	CCGCGTTGGG	CGTTTGACAC
23341	AGCACGCCCC	CGCCCTGCAC	CCGGCTCCCC	GAGGTCACCG	AAGGGTCAGG	AAACATAGTT
23401	TTCACGGTCG	CCAGTCGCGC	GCGTGGCAAC	ATCTTTCGGT	GCCCTCCGAA	CGACCCGGAA
23461	AAATCCCCGC	CGCGCCGGGT	CGTTCGGAGG	GCACCGAAAG	ATATCCCCAA	ACGCAACCTA
23521	AAGCATCATG	TTTGGGGTTT	CGGTGACGCG	CGCCGCAGAG	GAGGCCGGTG	GTGGCGCTGG
23581	CGAAGATAGT	GGCCGCAGAG	CACGGGTTC	GGTAGTGACG	CGCCTGGACG	CGCCTGACCT
23641	GGCAGCAGGC	CAAAAACACG	GGCGCCGCGA	GCGGCCGCGA	GGTGGCGCGG	CGCTTTCCCC
23701	CACGCCGCGA	TTTGGCAAAA	TGGCGGACCG	GCAGCGCCGG	TTGGACGCGG	CGGACGTCAA
23761	TCACGACGGA	GAGAAGCGCA	GTTGGTCAGT	TGTCCCGTCA	ATCACCAGAG	GCCGCGCCGG
23821	TTGGCGGATA	ATATGAAACC	GGCGCGGCCA	TTGGACGCTG	GCGGCAGCCA	ATGGGCGCGT
23881	TGGGTTTTGT	TTACAAGTTC	CCTATATATA	TTATATTATA	CCTTCCCTGT	TAAGGGAATC
23941	CCATGTTCTT	AAATCTAATC	AGCTGGGTTA	TATAATTAGA	ATATCAGTAA	TTACTATTAT
24001	TACTGTATTA	TTTATTAAAT	TATTTATTTT	ATTATTTATT	TAATAAATTA	ATTCTTGCGT
24061	TATTATGTAA	TCAGATACTG	GGAAATCTAC	TCCAGTTTA	ATAAAAATAA	ATAATTAATT
24121	CTATTATGGT	GGGGGTCTTA	AATATTGTAC	AAAACAACAA	TATTTTTTAA	TATTTTTCAC
24181	TAAAAATGCT	CCCATTTTAC	TTGTGACGTA	CATGTTAGGA	ATGTGGGAGT	GGTTTGTAAC
24241	TATGTTTTCA	AACAACACCC	TTTTGGGCGG	CTGAAACTC	ATATAAGCGG	AAGGCTCGGC
24301	GTTAATTGCC	ACTGGCCGCT	AACCAGTCCA	CCTTGCCAGT	TGGAGTTTTA	TTGCTGCTTA
24361	TGGCTTTACC	TCGCATTCCG	GGCACAAGCT	GATTTACGTA	GCACGCATTG	GGGGTTTAAC
24421	CCGGTTGAAT	TTAGTGGTTT	GCACGTGTAA	AAAAGCCAC	CGAGGCTTGG	GTTTAAGCGT
24481	TTAAAGCGTT	GCTGGTTTTA	TAACATGCGA	TTTAATTGCT	ATGGGAGCTA	GCTTGTATTC
24541	CGCATGCTTT	CGGTTTAGTA	GTTGTACAAA	ACAGAATTGT	AGCCCGCAAA	TTGTTACTGG
24601	TACCCAAAAC	GGCAAGCCCG	CCTAACACGC	CCCGCGCTGC	GGCTTTTGTT	GGTACCCGCG
24661	TCGCAATACA	ATCGAGATTT	TAATTTAAAA	TGGGGCAATG	GCGTAGCGCA	AATTTAAGCG
24721	CGTCAAAAAT	TTAAAATGGA	CTAACGTAAC	CGGTGCTTTT	AGTGACCTAT	GGCGAATTTT
24781	AAAATTTAAA	GCCGTGGGTT	TTAACACAGA	GCTGCCAGCT	TGTAATACGT	TGCATATACA
24841	GTGCTTTTGT	GATTTTATGG	TTAAGTGGAT	TTTAACATTG	AATTTTGTA	TGGTGTACAC
24901	GTGGATTTTT	AAATTGAATT	TAAAATTGTA	ATTACGTGCG	CGTGGATTTT	AAAATTGTAT
24961	TTATGTAACA	TTGTAAATT	TTAACTGCGT	TATGGTTGTA	TTTTGGTTAA	TTTATGTAAT
25021	TTTGCTTTTA	GATTTGTTTCG	GGTGTGAATT	CCAGACAGGT	AAGAGCCTAG	CTCCTAATGT
25081	TTGCCTTGCC	GCCTAGCTCC	TAATGTTTGC	CTTGCCGCCT	AGCTCCTAAT	GTTTGCCTTG
25141	CCGCCTAGCT	CCTAATGTTT	GCCTTGCCGC	CTAGCTCCTA	ATGTTTGCCT	TGCCGCCTAG
25201	CTCCTAATGT	TTGCCTTGCC	GCCTAGCTCC	TAATGTTTGC	CTTGCCGCCT	AGCTCCTAAT
25261	GTTTGCCTTG	CCGCCTAGCT	CCTAATGTTT	GCCTTGCCGC	CTAGCTCCTA	ATGTTTGCCT
25321	TGCCGCCTAG	CTCCTAATGT	TTGCCTTGCC	GCCTAGCTCC	TAATGTTTAA	CAACATTAAT
25381	GTTTAAGCAC	ACTAAAATTT	AAAGACGTTT	GTGTTGGTTT	TTATGACCAG	CTTGGTACAA
25441	AACCTGCTGG	TGATTTTTTA	CCCAACAAAT	AATAAATAAA	AAAGTTAAAA	CTTATTTCTG
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25561	GCTCCCCGTT	CCCCGAGGGT	CCCGGGCTCC	CCGTTCCCCG	AGGGTCCCCG	GCTCCCCGTT
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PCT/US99/26260

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81061	CTGAACGGGT	TGCCTGGATG	GCAGGTCTTT	ATACTCCTCC	TCATTGTTAG	TAAGCACCTT
81121	CATATCCCCG	TTCTCCAGGT	TCTTCACGCA	AATGCCAGTA	TCCGTAAGGG	TAATCAACAC
81181	ACCCTTCTCG	CACCTTTTGT	GCATAACCCT	TAGATCTTCT	CTGGCAAAC	GACAGTCAAC
81241	GACCCCGGGT	GACGGTAAAA	ACAAAGTCCT	CTCATCTTGG	AAACAACACG	CGTGGCCAGC
81301	GAGGACGGGA	CATGGTTCGAC	TGCAGACCCG	TATTCCAGCA	TGAACACGTT	CGAGGCCAAC
81361	GGTCTCTCCA	AAATATAGAA	ACGTTAATTG	GATGGCCCCC	GGCAGCCGAT	GCTCGCGCAC
81421	ACGCTTGTTT	TTGTCCATCC	GACTCCGAGC	CAATACCGTT	GCGGTAAAAA	CCCGTCTTCT
81481	TTCACGTCCA	GACCCACCCC	CCTGGTCGTA	TTGGAACCTA	CTGATTAAAC	CAAGGCACTG
81541	AGTTTCGGTG	GAAGACATCA	GATTACACAT	TGCACAACTT	ACCACCGGTT	CCGGCAACAG
81601	GCGAAACAGT	CTATATCGAG	CACCCGGTCG	CCCATAGGCC	TTATTAATAA	ATTTAAGCTC
81661	TCGTATACTT	TTGTGGTGTC	TCATGTTCGTA	CAAGAATCTC	ATTTTCCTGA	ATCTGCCATA
81721	GTGGGACGGT	TTTTCGCCCT	GACATATGCC	TCGCATGTTG	CAGTAGTCAT	CAAAAACTT
81781	CTCGCTATT	CATACGGACC	GATCTGCAGT	TACTTTGTTT	CACGGGACCC	GGACAACGGT
81841	CTTGTCTCTA	TTCTCCCAAA	CCAATCCGCG	GTACTCGCCC	GATTTCGACG	CGGCAACGAG
81901	CCAGGCCCTG	ATATCGACTC	CACGGCCCCG	CATTCCGCGT	ATTGGGAGCT	TGACGGATCA
81961	GCTCTCAAGA	CAAATGAAAC	CCTTGCGTTC	ACGGCACCAG	TTCATAAATG	TATCTAAATG
82021	ACGGAGTCAT	GTGCCCTAAA	CATATGGCTA	AGACTACTCA	AACTGTGCTT	AGGGGAAGAG
82081	TGACAAACAC	CTGGCACTCA	ATAAAATTTT	GAGGTCGGGC	ATTGACCCGT	GCCTTCCTCT
82141	GGTTAGTATT	AACCCCATCG	CCACCATATT	CAAGGAACAA	AACACTCCAG	ATACAAAACA
82201	CTTTATTACA	GATAGAAGGC	GCTCACCGGC	CTCACAAAAC	CGGTTCGCGC	CACAGGTAAC
82261	ATTAACATT	AGCGGCGGAC	AACGCCCGCC	ACAGACACTC	GTGCCACAGC	TGAATAATTA
82321	GCGGTACAGT	GCTTTGCACT	CCGCCCTCGG	GTGTAGGTAC	GCCCCCAAAA	TATAGGTAAA
82381	CACAAGCTTG	TGGAAGCGCC	GCTCCGTCCG	GAGATGTTCT	GGCCATGGCT	GTCATATAGT
82441	CAACCAAGTT	AAATGCACGA	CACGCGTTGG	CGCACGACAG	GACCGCAACC	GCCCCCGTAG
82501	ATTCGACGTT	CCCTGGAACG	AACGCCAGGG	AATGCCCCAA	CCATCTAATG	TAAATGCCGG
82561	AGGGGTTCGG	CCAAAACGCC	AACCCGCGCA	AAAGATCTTT	ATCCAGGAAG	CGCAACGCCT
82621	CGGCCTGAAA	CTCTGCTAAG	TGGCCTGGAA	TCTCCGGAAC	CAGTGCCTGC	TCGGCGATCG
82681	GTGCGGCGCA	TATGTGCCCT	TGAGGCGCAG	GTTCCGACAA	TGGGAGCACA	CGGATGCCGT
82741	TGGACGATTC	AGTCACCACT	TCACCAACTC	TGTCCCCATA	ATAGTAAAC	ATTATACGTA
82801	GCATCCAACA	CTGTTACCCC	CCGGCTCCCA	CCGACCAGT	GTACCGAACA	CCGGCGCCAA
82861	CCGGACCAAA	CTTATTCACG	GCCTCCTCGC	GCAACCCCCG	TAACATAGCA	GTAGTGCCGG
82921	AGGCCTGATC	GCAACGAGCA	CACGACCTCA	CGGTAATTGG	TAAAAGGCGA	ATGATTCCGG
82981	ATCTGTCCGG	ATATGGAGTC	GAAAGGTTGT	CCTTAGTTGA	CACGTCCGAC	ACGGTTTGAC
83041	TATGCCGTAT	GGCGCTGGAT	AGCCAGTTTT	TGCATTCTCT	AGCGGTTAAT	TCCCTGCCAG
83101	CCGCGTGCAA	GATGCCCTTG	GATCGACAGT	AGTCAATAAA	AATGTTTTTT	TCCGCTCCT
83161	CGACTCCTCT	GCTCCCTCTG	TCGTGATTCC	ACGGTATGCG	GATTAAAGTT	CTTTCCTCGT
83221	CAACCCACCG	CATTCCGGGA	TGACGACCTG	TCTCGCAGCA	TTCCACCAGC	CATCCTCGAA
83281	GGCTGGATGG	TTTGGTCACT	CGGACCGGCC	GCTCCATATC	CTCGCCGCGT	GCACGGCAAT
83341	GCCAAACCTT	GTCCAAGTTA	CCTCCTGACA	AGCACCTGCG	ACCCTCACAG	ACCTACGCGT
83401	GGCAAACGGG	AACTTGTTGG	CTAGTCTGCT	GGGACAGTAC	CTTAGCGTTA	TTTTTATCTG
83461	TCGTGCACTC	CTGCGGCGGA	AGTTTGTGGT	TTATGAGCAC	CACCGGGAAG	GATTTCATGTG
83521	TCCCTGCACC	AGCTCGTCCT	GACAGACCAC	CACAGAAATA	GGAACCTTGT	GGCTCTCTTG
83581	GGCACCAGGC	TGCTGTGAGA	GATACAGTGT	TGCCCCGTTG	CACGGCGCAT	GGCTCTCTGG
83641	TCCCGGAGAG	CGGGCAAGCT	TTCTCATGAA	ACCCATGAAG	TTAAAAATTT	TCTGTTTAGA
83701	GAGGAACATG	ACCTCTCGTT	GGATAATATC	ACCATTACAG	TCGTTACCCG	CCGCGTACAG
83761	CGTTTCCCTG	TTCCTATTAT	CGACGTACAC	TCCGGTTTCG	TTGCCGTGTA	CTGATGCC
83821	CTCGTTATAC	GCGACCAGCA	TCGTTTTTAT	TTGCGTTATC	TGTTCTGCGG	TCAGTTTTTC
83881	ACGATCTGGA	GATGGAAACC	AGACAGTGAA	CCCGGTGCCA	TAGAAACACA	GATGACCGGG
83941	ATGGGGAAAC	GGGCGCGGAT	GCAAACGAAC	ACCTCTCCGA	TCTTGGGACG	TGTGATCCAG
84001	TGCCAGAACG	CCAAAATAAA	ACACCTGAAT	TCTCAACCGC	CAAAACGGCA	CAATTTGATG
84061	AGGCAGCGGC	GCTTCTGGGA	CCCCGAGACG	ATGTGCTCGG	GCGTACGCTG	ATCTCGTCGG



84121	AGGGACCAAC	GTGGGCAAGG	GGGCGGGCAG	CGGCATTCCC	GGCTCTAACC	CGATGCCACA
84181	CACCTGCTCC	AAATAATTTA	TAATCCCTTC	AGAGTTCTCA	TCAAGGATAC	ACGCATAGCA
84241	TCCAGAATCC	ATCAAAGGTT	TTACCCGGA	GGCGGTCCAG	CCGCCACCGT	GCGTCCGCAC
84301	GACCCCATGC	TCCACTTCTT	GAAATCCAGC	ATTCTCCCTC	AAACCCCTCA	AAAATCGTCT
84361	CTTACACTCT	AGCAGGTTTC	CATCTAGCCC	AACCTGCAGA	CCTCTCTCCA	CGCAGTAGGC
84421	GACAACGGCT	GGAGTTCCAG	CTGCCCCCTT	CTTCCAAGAC	AGCTTAAAGC	TTTTTTTCCC
84481	TCATCGCACC	AAGTTAGGTC	TGAATAGGTT	TTCTCGTTGA	GATGGTAAAG	GGTCCACGAT
84541	CGTAGATGGC	TGTAGTTAGA	ATCTCGATTG	GCCATGACAG	CGTACACCTA	TACAATAAGG
84601	CGCGATACGA	GTGTACTTTT	CCCCTGATGA	TATTTGCAGG	GCTCTATCTC	CTCAAAACGT
84661	GCACCTGGCT	AAGCAAGCAG	CTCTATTTCT	GCCACTTCCT	CATTTATAAA	ACCATATTGT
84721	TTCTGGTCTC	ACGTTTTGCA	GCTCACCTGC	AACGACACGG	TGAATGATCC	CGAGTTTCGC
84781	TCCTCCAGTG	CCTTGACAAA	CACTGGCCCA	GAGTTAAAAA	GTGAAGCAAA	AGGCATAGCT
84841	TTCGATTTC	GTCACCGGCA	GTGAGTGGTG	GACATTAGAA	AGTGTGTTGGC	CCACACATTC
84901	AGCTGTGCAC	TCAAGCCCGA	CCAAATATGGC	ACGTGACCAT	TCCCCTATCA	CAACACTAGT
84961	ACAGAAAAAA	CGAAACCACA	CGCGAGGTGA	CATTTTCCAG	GTTAGAGAAA	TTTATTAAGC
85021	CGGCGAATCC	ATCTAACAAA	ACAGCAAATG	TTTATTCAAA	GTGCCTATAG	ATTTCTGTTT
85081	GATAAACAAAT	AATAGATAGT	GGAGCCCTCG	AAGAGTTTCG	CTCCTCGCCC	AGGGAACAAC
85141	CCGAAACCCA	GAGTTTAACA	AAGGCCGCGG	GGGGACCCGT	GTCGCGCGGG	GAGGGGGACC
85201	TGGCCATATC	AAGCAAATAA	CGATCAGTGT	CAAATGCCCA	CACGTGCATC	CCGGTGTAGG
85261	CACGATGTGG	CAAACGTGAC	GGTCCACGG	TATTTCCCTG	AAACCACACG	TTAGGACCCC
85321	CGGTAATCGT	GAACATAATT	CCCCTGCTGG	TGCTGCAGAA	AATTATTCCA	TCCACGAGCC
85381	ATTTCAAAGC	GTCTCGAGT	CTCTGCACCA	CAGACGACTC	AGTCGTCTGT	GGTATCGGAG
85441	GAAACCACGC	CTGAAGGGGT	CCATCCAAAA	CACACACGTG	GTGTCCCGCC	TTATTTGTTG
85501	GTTTGTGAAGA	TAGCCTGATT	CCCTGCCCGG	TGGCTGTCAA	CAGCTCAGCC	TGCAATTGCC
85561	CATAGTAATA	AATCTTTATA	TGCATGTGAC	CCCATGGATC	GCCGGCAGAT	GGCCGACAC
85621	ACGCCATAAG	TCTGGCGCGC	AGAAGGCCCG	AAGAATACGC	TGCATCGTCG	TCCCTGCGCGT
85681	CGTGTCCATC	TTGATGACCT	TCCCGATACC	GTCTCTCTTG	ATCACTTACG	GACGGTCCCG
85741	AAGGGCGCTG	AAGCATTCCG	CGCCCGAGCA	TGGCAAACCG	CATATTGCGC	AACTCAAGGC
85801	GGAGCGCGCA	CACGCGCGGA	CACACCGGAC	AAGACATCTC	TTCGGATGAA	CGCAACCTAA
85861	AGACCAGCGC	CGTGAACGGA	AATTGATCGG	TAGGGAATTC	CTTATCTTCC	TCAAAGTACT
85921	TGCTCTTTCT	TACGGCTCCC	AGCAGGCGCC	CCCTGGCTTG	ACACAGCCTG	CTCCTTCCAG
85981	ACGGCAGTGG	GATGTGTCTT	AACTCACAGT	AGGCATCATA	GTGTTGACCG	TCGCGTTGAT
86041	AATCATAATT	GGGAAACGAC	GGTGGTGTAG	CTGCCAACAC	CAGCCTCGTC	TTTTGCTCGT
86101	CTGCCCAAAA	CAGGCCCGGG	TACAGACCAC	TATCGGCTTT	GAACGTCAGC	CATTCTTTGA
86161	GGGCCATGCC	ATAACGAGAT	GGGCCCTTTC	TGCATTCCCC	AGCGGCCATG	ATTCAAGTCG
86221	CAACGTTTAA	AACCCATGTA	AAGTTTCAGT	TCAATTTAAA	ATGACACAAC	TCCGCCCATG
86281	CTGTGACAAA	CATAAAGAAA	GTGGCACACG	TGTGCGCTTT	GGCCGTCCTG	TCAGATGAAC
86341	CCCGCCGGTA	ACATTTACTT	CCTCCCATAC	GGTGCAGAGG	TAGATACAGA	TGTGCTTCCC
86401	GTTGTTCCCG	AAGCACCGAC	ACCCGCTCCA	GACGGTCCCT	CGTCCCCGCT	GTCATCTCCC
86461	GCGGCACCGT	TGCTGACATC	ATCTGACACA	TCATCATTGC	ACACGTAGCG	AAGTTTCAGG
86521	GCAATGGGAG	CATCCCATGT	ATTTTCGGAC	TGAGTAGACG	GGCACATGAA	AATCCACGCA
86581	CAGTCACGCG	GCGGGACATC	TCCTGGAGTT	GGCGATCTCG	CCAGCTCCCT	CAAATAATGC
86641	AAGAGATCGA	ACGCCTGGAG	CGGCTGCGAA	GGCCGAGCT	CATAGTAATT	TTCAGAATAG
86701	TTGGTCAATG	CCGTATGTT	GCGTCTTCC	AGGTTTTTCA	CACAGATTCC	AGACTCCGTC
86761	ATAACACAGA	TCAAACCTCT	CTCGCATGTT	TTCTGCAGGG	CCTGCAAATC	TGCCCTGGCA
86821	AACTCGCAGG	GAATCAACTG	GGACGACGGC	AAGAAACACG	TCAATGGATT	ATCGGCACAA
86881	CACAGGTGTC	CGGGACGTTT	GGGATCAGGA	AGGCCACATA	TACGCACACC	CGAGCCGGCT
86941	CGCAAAATTT	GAACCTCCGAC	GTTCTCTCCA	AAGTAAAAAA	AGGATACCTC	AACAGCCCCCT
87001	AGCACAGGTT	GCGCAGAAGA	GTCCTGGAGT	CGTTTGGATT	TCCGTTTATA	AACGGAGACT
87061	CTCAGTGGCG	CCTTAGCCCT	CGGCCGCCGT	GTCCTCTCGA	AAACATCCTC	TGCAACGCAA
87121	CTGCCCATGG	AACAGCTATG	CAACGATGTG	GTCGCGATGA	GGTTACACAG	ATCCGAGCCC
87181	ACCGTAACCT	CAGGAAGCAA	TTGAAAGAGT	CTGTATCTTT	CCCCAGAGAC	CCCCCCGGCC
87241	TTGCTCGGCG	GTGTCAGCTC	CCTCAAATAC	CTGTGAGATC	TCACTGCATA	TAACATCCTA
87301	ATTTTCTTAA	ATTTCTGGGC	ATGTGAACCG	GTTTGACACA	CGCCACGCGC	TGAGCAAAAA
87361	TCGTCAAATA	TCTTGTTATA	ATCTTCATCG	ACGCGACTGT	CGGTGCACCG	GTTCCACGGG
87421	ACACGTATAA	TGGTCTTGTC	CTCGTCATCC	CATTCCACTC	CGGGGTATCG	CTTAGACTCC
87481	ACGGCCTCGA	TGAACCAAGC	TTTTAAATCC	ATATCGCGTT	CCGCCATCTC	GCGTATTCAA

87541	TACCTCTGAC	CGAGCGGCTC	AGTTAAAATA	CCGAGCCCCC	AGAACCCAG	GCCTCCGTAA
87601	CAAAAAGGAT	CTAGATCTGA	AAGAAGCCTA	ACTATATACG	CCCACGGGAA	AACCTCTGTAA
87661	CAAATGAAAA	AAATGTTCTC	AGAAAACGAA	GAGAGGAAAA	CTGACGACAC	GCAATCTAGT
87721	CAAACCAGAA	CCTAAAGGCC	GTCAGTGACC	CATATCTTTC	CCCTTCCCGA	GCGAGTGCGG
87781	TTTCTCACAT	GCCGTGAGCA	AACGAAAACG	CTACACATAA	TAAGACACGT	GTGAGAGGAA
87841	AACTTTATTG	CAGGGACAGG	GCAAAAGCAA	GCTGTGCACG	GTAACAGTAT	GTGTCACTGG
87901	GCCGCATCCC	CGCACGCCCC	CCACAGACAC	TCGTGCCAAA	TTTGAATGGA	CAGGGGGATT
87961	GTGGGACACG	GCTCGCCTTC	CGGAGTGGGA	ACGCCTCCTA	GGTGGAGAGA	CGCATAGGCG
88021	GCTGGAAATC	GCGTTCCGTG	TGGCGGGGTG	GCTGACACCG	CACTCAGATA	TTGGGCGAGC
88081	GAAAAACACG	GCAGAACTCG	CGCGTTGTTA	ATTATATCCA	CCGGCTCCAG	CGAACCTCCG
88141	TTACCCAGCA	GGAAGGCTGG	AACGTGACCC	ATATACCGAA	TGTAAATCCC	AGACCCGTCT
88201	GCGTAAATCA	TCAACCCGCG	TATGATCTCC	TTCCCCAGCC	ACTTTAGCAT	GCTCGTCTGG
88261	TGAGGAAACA	TATCCTGTGG	GGTATGCGGC	AACAGGGCCT	GTTCCGGCCG	AATGCCCCGA
88321	CAAACGTGTC	CTTGCACTGG	TCGCTTTGGC	GCCGGCAACA	GGCGAACGCC	GTTGGGCGAC
88381	CCGACCTCCA	TGTTCCCAAC	CAACCGCCCA	TAGTAATAAA	CAACCAGCCG	TAGCATCCAA
88441	CACTGCTCCC	CGTCGCCTCC	CACGGCGCCT	TCGTACTCCA	TCCCGGCACC	GAGATACCCG
88501	AATCTGGCGG	TTACCTCGTT	GCGTACGTCC	AGGAGCATGC	CGCCGTGGC	ATCCGCTTCG
88561	TTACAGAGCG	GGCAAGATCG	TACAAGATC	GGCAACAGGC	GAATAACGCG	GCACCTGTTA
88621	GGCGCCGGCG	CCGACAGGTT	CTCCTCAGTG	GATACATCTT	CGACGGTCTG	ACTGTGGCGT
88681	ATAGCGCTGG	CCAACCACTT	CTTACATTCC	CTGACGCTCA	ATCGTCTGTT	ACCGGTCTGC
88741	CGTATGCCTT	TAAATTGACA	ATAGTCCAGA	TACATGTTTC	TCTCATCGTC	AGAGACGCCA
88801	CCCGCGCCCT	TTAAATGATT	CCAAGGTAAA	CGAACCAGAG	TCTTTTCCTC	ATCCAGCCAA
88861	TGCATTCCAG	GATGCTTATC	ATTATCGCAA	CAATCGAGCA	ACCATGCCCT	CAGGCCAGAC
88921	GGCCGGTTTA	CGCGGATCGA	CCCTGCCCGT	CCTTCCGCCA	TCTCTGAGGG	TCCGCTCGCA
88981	TAAACTGGCT	TAAAAATCTA	TCCCCCCCCG	TGCACCTGTA	CTTTACAGTC	ACACCTTCGG
89041	GCACACGCCG	TAGGGCTGGC	AGGGACGGGA	CCCAGCAGCT	ATTTTCATCT	ACACGCCGCG
89101	CTACTAGCGG	AAATCTGCGG	TCTACAAAT	CCACCGAGCC	GCCCCAATGT	CACCATGAGT
89161	CAATTACATC	TGACAGATCA	CCAAAGTGAC	AGGGACCTGG	GGTCTTTTAC	TGGAGCCAGG
89221	GCGACCCCCA	AGGTAAAGAG	TGACTGCGTT	GCTGGGAACA	GTGTGCGCCG	GAACGGGTGA
89281	CCGGGCGAGT	CTGCGCAGTA	GACCCATAAC	ATAGAAGATT	TGAGATTTGA	CAGAGGATCG
89341	CACAGCCCGT	TGTGCGTGGT	TCCCCTCTGC	GTCGTTCCCT	GCGCAGCGAA	CCGTGTGCCCT
89401	GGTTCTGTTA	TCCACATACA	TCCCCAGCTC	ATTGCCATGC	AAAAAGACGC	CCTCCTCACA
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89521	AGGTTTGGGA	AGCCACATCT	TATACGCTGT	CCCATAGAAA	CACTCGTGAC	CCGGTTTGGG
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89641	CGACCCAAAA	TAAAAAACTC	GTATACGTAG	CCGCCAAAAT	GGAGTGATCT	GGTCTGGATG
89701	CGGTGGTGCA	ACCGTCGCCA	ACCTGGCGGC	CCTGGCATAT	ACGGACCGCG	CCGCACCGTC
89761	AGCCTCCGCC	GGAGCGGGAG	GAGGCAGGGG	CGTCCCTGGC	TCAAGCCCAT	GGACGCAGGC
89821	CGCCTCTAAG	TAATTTAGGA	CGGTCTCCGA	ATGACTATCT	AAGATGCAGG	CATAACACAG
89881	CGGATCCACG	GCGGGTCTCA	AGTAAAAAAA	TCTTTCTCCG	CCAAAACGCG	TTGTTGTTGC
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90001	GCACTCCGAC	ACGTTGTGCT	GGCGTCCACA	TTCCAGATCC	CTATCCAGGC	AGTACGCAAC
90061	CACAGGTTGC	ATGCCACTCA	TACCTCGGTG	CCATGATATC	CTAAAGCTCC	TCTTTTCCTC
90121	GTCGCACCAA	CACAAGTGTT	CGGGAACGTT	GTTTCGCTTC	AAATTGGAAA	TAATCCATCT
90181	TCGCAAGTGA	TTGTGAGTGA	TTTCTATCTC	GGTCATTGTG	AACTTGGTCC	ACGAAAACTA
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90361	CGTTTCGGGT	CGCTCACCTA	AACTAACTCG	AGCATTACAC	GGTTCATAGT	GTCCCGGGTT
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90481	ACTTTATTTA	AGGAGAAAAA	AACCACCGCC	ATCAGAAGGT	TTCCGCAGGC	ACACACCCCC
90541	AAGACTATTC	TTGGCGAGTT	CGCGGCACGC	TGTAGTTTTT	TTCTGATTAT	TTGTAATTCTG
90601	CATCCAATCG	CCCCAAACAT	AGTCAATAAA	AAAACCGTAA	ACACACACAC	TCTGGTGAAC
90661	AAATTATGCG	TTCCTGGCCC	CAAGACGTGC	GTGAGCGTCA	GCAGCACAGC	GCCTGCTAAA
90721	ACATAAACCA	GGGCTAAAAC	GTTTGACCGA	AACACGAGTC	CGAGGCAACA	AAAAAGCCCCA
90781	ACGTGGAAGA	CAAAAAAATA	GTGTACAAGT	CCGAGGAGCA	GAGGCGTCGT	CAGGCCAACA
90841	GACAGCGTAC	CCGCGGCCGC	CCCTCCAATC	AAAACAGTCA	CGAGATAAAA	GTCCCATCCA
90901	CACACCTCAA	GCGCTTGCTT	TATAGAGTAT	GTGATGTACC	GCCTAGAGGT	GAGCGAAAAA

90961 ATTGCATGGC GTCGCTGCAG GCCCCCCTCA TAAAGCGGAT GTGACCGCAG CGCTCTGAGT  
 91021 GAACAAACGC CACCTACTAT AAAGGCCACC AGGGGCAGCA AAAACACCGG GGCTAAAAAA  
 91081 AGGTTCTTAA GAACTAAAAA ATAATACACC GAGAACGCTG CTAAAAATCC AAGTTCGTAG  
 91141 TATCTGTGGG CAATCGTGGG GCACAAATAT ACCACGTCAT TCGCCGCAAC AACACAGAGC  
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 91441 TACAGCGAGT GATAAATAAA ACACAGAATA AACGGGGAAG CCGAGACCGT CCCGCTCATA  
 91501 AAAGATAGCC AGGCGAGGGA CGCCTCCGAC GTGTACGTGC CCATGGCGCG GCGTTTAAAA  
 91561 CAACGGGTTT AACGTCGCGC GAGGCGTCTT GGTTTTTTCCT TTCCCGCGAC TGGGTGAGCT  
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 93901 ACTGCAGGGC ATTAAGTTTT GTCTGTTTCT GAATTCGACA ATAATACAGC ATGGTTTTTA  
 93961 ACCCTAGCCT GTATCCGGTC AGGAGCAAGT CTCGAATATA ACTCGCCCGC GTGGCCTGCT  
 94021 CCTCCTTTAA AAAGAAATTA AGCGACTGAC TCTGGTCTAC AAATGGAGCC CGCGCACCGG  
 94081 CCCGCTCCAA CTGCTTAATT GGGCAGTAAT CAAAGGCCGT CAAAAAATC TTGTATCTGT  
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 94201 CACCAGGCTT CACTCTCTTA AAAAAAGTCA CGTTTGGCTT CAAGATCTCT TCTTTGCTGG  
 94261 TGACCTTCGA TGCGATATTG CAAAGAACG GATAAAATGC TTCGGTGATC CCCGTAAGCT  
 94321 GAGAGGTTCC GGCCGTTGGC ATCAGCGCCA AAAACTGGCT GTTGAAAATT CCATGCTGGG

94381	CAATGCTGCG	CCCCAGCTGC	TCCCATCTCT	CCAAGGGAGG	GTGGGACGGC	TTAACGCCGT
94441	CCCACGTTTG	CCAGTGAAAC	ACACCCTGAG	CCAATCGACT	CCGCTCCCAT	CCACGAAACG
94501	GAGTCCCCTC	GCCGAGTAAA	ACAATCTCAT	GACTGGTTTG	CACCGCCGTA	AAATACATTG
94561	CCTGAAAAAT	CTCCACGTCC	AGCTTGGCGC	TCTCGGCGTC	AAGGTAGCCA	AAGCCCAGTT
94621	CGGCAAACAC	GTCGGCCAGT	CCTTGGACGC	CAATGCCCAT	TGATCGTTCC	TCCTGGCCGC
94681	GCCTAACGCT	CTCGGTAGGC	GCCGTCCCGC	CCAAAATGCA	CGCATTGACA	ATGATAACTG
94741	CGGCCTCCAC	GGCATCATCG	AGCAGTTCAA	AACCAAATGT	TACATCCCCC	TTCCCAGGCT
94801	CCACACCAGA	CTCTCCCCTG	TGGGGTCTGA	GGCACTTTGG	TAGGCTAATG	TTTGCCAAGT
94861	TACACACGGA	AGCCTGACCC	TCGGGTTGCT	GCACGATTTT	CGCACAGAGA	TTAGAACAGT
94921	TTATGGCGCT	GCCCTGCGTC	TCACACCAGT	GGTGTGTTGT	GAGCGCCTCC	TTTAGCAGGA
94981	CGTAGGGACT	GCCGGTCTTA	ATGACAGTGT	TAATAAGGGC	ATACATCATC	GATTTTAAACG
95041	GCAACGAGCT	AGAGTGTTTG	CCAGCGGCCA	CTAGTCTGTT	GTATTCAATC	TCAAATTCGG
95101	CACCGTAGAG	TTTTAGGAGA	TTTGGCGCCA	CCTCTGGCGC	AAACAGGTGC	CACTGGCCAT
95161	CTGGGTTTGT	TTCGTACAGT	CGGAAAAAAA	GCTCCGGCAC	ACACACGCCC	TGAAACAGGT
95221	TGTGACACCG	CTCCTGATTC	TCCGGCATCT	TCGCGTTCAA	AAAATCACAA	ATCTGATGAT
95281	GCCATAGTTC	CATGTAGGCG	CTGGCACCAC	CAGGACGGAT	ATTGTTATCC	TTGAAATACC
95341	CAACGTGGGC	GTTTATTAAT	TTTAAACAGC	TGGTGATGTT	CTTGTGTTCC	GCAAAAGACG
95401	AGACATCTAT	CCCCACGCCT	GACTTGCTAG	CGAGAAGGGG	GGACATTTCC	TCATGAAGTG
95461	CTTTGAGGGT	TTTGTCTTCG	GTCGCCATGG	ACGGCTTTAA	AATAAAACAG	CTAGAAAGCT
95521	GACCTCCGCG	AAGCCCAGGT	GACCTTAATA	CAGGCGTTGC	GCAGCACACA	ATCTGTGACG
95581	AGATGTAATG	GAACGCGTAA	CCAACCAGGT	ACATCTCATC	CAGCTCCGTT	TCGCTCTCCA
95641	CCAGGTGTCT	GAGGGTCTCC	CGCAAACACG	GAAATTTTAT	AACTGACAG	GCCACAAAAA
95701	CAGCCACCCT	CATAAACATC	TGGGCCACGC	TTTCAAAAAT	GGGTGAAGAA	CCCTGGGTTT
95761	TCAGCACGTA	CGTATCGTAA	AACCTAACGG	CAGACAGGTA	GCCGCAGTTA	ACGAAATTTG
95821	TGTATGCCTT	GCTCTGCTTA	AAGTCTGTGA	AAAGACCGTC	AAGGGCCGCT	TCGTGTTTTG
95881	ACATAAACCG	GCGAACCTCG	TCGCTTAGCC	TTTCCCGGAA	TACCGCGAGA	TAGTCTCGCA
95941	CCGTAACCG	GCACCTGTCC	TCCATTATTC	TGTGCCAAAG	GAGACCGGTT	AGTGAGTTGG
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96061	CAGGGTCAAG	GCTTGAGCTA	CACCCCCCAG	TTCCCGCGTC	GGTAACTAGG	GTTAAAGGTT
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96181	AGACCCCGGA	TTGGAACAA	ACTCCGTAAA	TTTTAACACC	GGTAAAGCAG	CGCCTTTAAA
96241	GTGAAGGCTT	TGAAAAGATG	GTTGTAAACC	GGAAGGCACG	CTTCCAAGTC	TGCAAACTAC
96301	GCCGAACGCA	AGCCTATTTA	TATACAGGTC	ATTCTGCAGC	TGAATGTATT	TGGTGCGAAT
96361	CACGCCGCTG	TAAAAATCCC	TCAATTGGGC	AGCTATTTCA	CAATATCCTT	TACCAGACTT
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96481	AATAGACGAG	TGGGCGGGCA	AAGGCTCGTC	GCGCAGGCTG	GACGGGCATG	AGTCATCTAA
96541	ATCCACAAAC	ATGTCACTAG	GAAGCGTAAG	GCCAATATGT	GTAACAACGG	GCTCTCTGGC
96601	GACTACGTTG	CCCTTTAACG	CAGACGTCAC	CTTGGTGACA	AACGTACTGT	GGACCGTTTG
96661	AACCAACGGC	CCGACCGGCG	CAAGAACTG	ATGAAGCGAG	CCGGTTTCCA	ACAATTCTTC
96721	AAAATTGGGT	ATGGCGTCAA	GTAGACCGCT	CTCGTGGCCG	TACCAAACAC	ACGCTATTCT
96781	GTTGGTCTGG	GGGGCAGAGT	CCGCGTCCAT	CCTAGACAGT	CGCGCCAGCG	ACGTAGGCGT
96841	GAATAACATG	TCAATGGAGG	ACCCAGTGTC	AGTCTGTTTA	AAGGAAAACA	GGTAGGTGCC
96901	CCGAGGTTCC	TGTGAACTCA	TGGTCTGAGA	ATAAATCAAA	AAATCTCCAT	ACGTTTGACA
96961	TGTAGGCGAG	ACAGATAAAA	ATCCATCTTT	GATGGCCTCC	ACCCAGTGG	TGGTCGACAC
97021	CACATATTTA	GAGAGCAGAT	CACGAACACC	CTTAGAAAAG	TCGCGACCGC	GAGATAACGA
97081	AACCCGGTGA	GGAGGCGGCG	GCAGTAGACG	CATCAAATA	TCATTACAGT	TGTTACAGTT
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97201	GAACACCGGC	GGGCTTTACA	AATTAACCTGA	CAACCTCCTA	ACGTGCACCG	GATCGCTACA
97261	ACAGCTTAAA	CTCCTGATGG	AGTTCCAAC	AAAACCACTA	CCAACCGCAC	ACCTTTTAAAG
97321	CATGCCACCC	GTGACCCGGT	TTTTAAATAC	TGCATTCAAA	ATAGACAACC	CCCTGGTTTC
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97441	CATTACAGAC	CACCAAAGCG	CAGAAACGTC	AACAGGCATA	CTATCCGAGG	TTGTGAATGT
97501	TCTTAATACA	GCTATTCGAA	AACCAACGGA	GTCCCCAGCG	GCTAAAGACA	ACGATTATCT
97561	CGACAACCGT	GCCATATTGG	CCATGATTAC	AGAATACATC	CATCACGTAA	CTTCACGTAC
97621	GCCCTCGGGG	ATCCCACCGA	CACCACCAAT	GGGTATCAGC	CATCTACCGT	GCGTAGAGCA
97681	AATTTTACAC	GAAACCCACC	GGCAATACTG	GAACCTAACC	CTCCCGGAGT	CGCTATTTAT
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101641	AAAATTCAAA	CAGGGGGCGT	TCAAAAAACA	CCTAACAACT	AAAATCAACC	GGTGTCTGGC
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101761	AGGCTCACAG	ATTTTCGTCCA	GCGAAGCGAA	AATATCAGTC	CGGGCGCTGA	AGGAACAGAT
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101881	ACGGGATAAA	ATTCAGGACC	TAAAAACAGG	CATCGAGCAG	CGCAACAAAG	AAATCCAACA
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103561	AGCGTTTGAC	TGGATCGCGC	CCTATCAAAC	ACGCGTAAAC	GCGTTTCTAA	AAACCATAGG
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103681	CGCGGTGCAA	TCCGCAGACC	TTCAACAGGC	CACGGTGGGA	ACAAGTTTAG	AACGACCCGC
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104161	TTTGAGTTCA	ACGGTATACG	ACAAAATCCT	GGACAAGGAG	CCTTACGAGA	CAGCCATAGC
104221	GGGATTGCGG	TGGCTGGAAA	TCGCGACAAA	ATCCGTAATG	GTCTACAGTC	AACAAAACGA
104281	AACGCAACAG	TTAAACGTAC	TGCTGAGCGA	GGTAGAAAAA	CAGAGCACCG	TCGCGCAGCG
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104401	GCTAGACGAA	CTCGGCCCCC	TACGGGTAAA	GGCGGAAAAA	ACCACCGTAG	ACGCGTGGA
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105841	GCCGGTCCAG	GCTTCAAACA	CACCGGTATC	CGCTTTCGAG	GCGGTGCTCG	GAGCAATGGT
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106321	GTCCGTGACG	CACCTCAGCC	ACGACGAGAT	ATGTAACCTC	TTTACTACGC	TATCCCGAGA
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106741	AACGGCCCAT	ATAAACCCAG	CAGCCTCCGG	CGAGGTCACC	GAACCAAAGG	GAATCTTTGG
106801	GACGTATAAA	CCCCGAGTGC	TCACCGAACC	CGCCAAACCC	GCAAACGCCG	GCGTAGCCTC
106861	TCGCCAACCA	GAGGCAACCA	CCACGGTCCC	CAAGTTACCG	ATTAATCCAC	CCACCGCTAG
106921	GGTCTTTATA	GGGACCGCGT	CCAAACTCTC	GCCAGCCGTC	GAAGAGAGCC	ACGGCGCCAC
106981	ACCCGACGCA	CATCAGTCGA	AGATAGATCG	GGAAAAATAC	GCCGAGAGTC	GGCCTCGCCG
107041	CACCCACAC	CTCGAAGAGG	GGCCACGGGA	GCCTCACGTC	AACACTCCAA	CCAGCGCACA
107101	CATAAACGTC	CCCTCTAGCC	AAGGTCAAAA	AACAGTACAC	GGGCGCGAAA	ATCCCGGCCCT
107161	TCAAACAGCA	ACTCCCAGCG	CCCCCAAC	AACCGCATCA	AACCCGCGCA	TTCAATACAC
107221	GCTCCCCAGA	ACGGACGGCC	GGTTGCTTCA	CGACGAATCG	GAGGTGGAAT	CGACCCCAAC
107281	CGAGGAGGTA	AAACGATCGC	CAAAAACACA	AGATGTGTCT	CACGGGCCCC	AACCGGACGA
107341	TCCAGGTGG	ACCGCCCCGC	TCGGTCCAAC	CATAGAGATT	CATCGACTGG	AACACCCCCA
107401	AATTCTCAA	AATATAACAT	CACTCACCGT	CCCCACTCCC	AGAGTCACCC	CAATCCCTCC
107461	CACTAACATC	TGGATACCCC	TATCCCACGT	CAACATCCAA	CACGAAGAAA	TCACACGAGC
107521	CAAGAATGTG	TTAATGCGAT	TTATTCAAAA	CGTACGAAGA	AAACTTCAAG	CGTCGTCTGA
107581	CGCTCTATCC	GAGGCTATTG	CCAGAATAAA	GTTTTTATAT	CTGTAACGCG	CCCATCTCAC
107641	TTGCTTTTTT	TATTTTGAGA	CGAGCGTCTT	GTGTCCAGAG	TAGTTGCGTC	GCTAGGTGAT
107701	AACGAAGTGG	ACCCTATGCC	AGAAGACGCC	ACGCTCCCGG	GTCCGCCGCG	ACCGGGCGCG
107761	GGGCCCATCG	AGCCTATCAA	TGAATGGGGC	CCGCTGGAGA	TCGTAGTAAA	GCTATTTGAC
107821	CCGAGGGTGG	AGGCCACCGG	GGCAACGCTC	GGGCGGCGCG	AGCCGGACCC	GGACAAAACG
107881	CCCAGATAC	TAGAACTAGC	GTCTTTTGTG	CCGCGAAGGC	CCCGGAGGTG	GTCTTTGCGC
107941	AGAATTCCAT	TCTTCTTTTG	CATATACATG	TCGTAATGAT	GTTTGCCCGT	TAAAAACACC
108001	AGATAATTAC	GTTTCGCGAT	GGCATACTGG	GCGGGAGACA	TGTCACCCTG	GGGAAGGTTG



108061	TTCATCTCGG	CAACCAGCGG	GTGATTTGGG	TAATCGTGCT	CAAGGCGTCC	CTGAACGATT
108121	GGCTCCTTAA	CCCACAAGGA	CGACATCTCT	TTTAACTTCT	ATTACACTTT	CCACAGGACA
108181	GGGACGATAT	AGACGAGGTC	AAATAAAACA	GCTCGGCCAC	ACGCAAATGC	TTTAATAGGC
108241	CGGTCGCGCA	GTCCGGCGAC	GCCAAACAGG	GCACGACGAC	GCTAACCAGG	GAGTTCGCGT
108301	CTCGTATGGC	ATGTGCCGCG	TTGTTGGCGA	GCACGCACCG	TAAATAGGGA	TCTCCAACAC
108361	ACGTGATCTC	GAATAGAGAT	ATAACCCGCA	TGTGCGATCC	GCCACAATAA	GAGCAATATA
108421	CGCGCCCGGT	GGTAGCACAG	ATCGAAAGCT	GCTTCTCTTT	TTGGTCGCGA	CTGAAAAACA
108481	CGTTGGTGGG	TGGGAAATTT	ACGGTTTCAA	ATTTACCCCG	TCCGAAATTC	AAACAGTAAC
108541	CGCACTCGAG	GCACACCACC	ACCTTCGGAG	CTGGCACGGT	CTTCTCCAGT	ACGCTCCTGG
108601	CCACCACCTG	GGACCAAACA	GGTAGAGAGA	TACACGGAAA	CAGTACGTTA	TACGCCAATA
108661	CTTTTTGACC	CAGGTCGCGG	GATATCTCCG	TCTCGGTCGA	CTCCCCTATG	GGCAACACAA
108721	CACGGGACAT	GCTCAGCAGG	GCCCTAAACG	TCAGGCTCCT	CAGAAGGGCG	TTAAACGGGT
108781	TGCCGCACGG	GACGGTCGGC	GCCAGTTCTC	GCAGCGAGGC	CAGAAGTCCC	GCGTCCGAAG
108841	GGCCCGGGAC	ACTCTCATTC	AGGTTAGCTC	CCAGACGTCT	GGAAATGGAC	GGAACGTTCA
108901	ACTGCATCGA	GACACAACCG	CCCCCGTTCC	ATTTCTTCCG	CAAACGGGGC	AGATCCAACG
108961	CGTGCTGTGG	CAAACAGGTA	ACCAGGGGAA	ACCGCTGGCG	ACAGTTAAGG	GTTTTGCACA
109021	CGAGACAACA	CGCCCTCTGA	AACGACACAA	CAAGTACCTT	GGACCACGCG	CTCCTGGGAA
109081	CGGCCGTTAA	ACTTAAACCT	TCGTCGCTGG	GACAGCCAC	GCCGGTGCAG	ATACACCTCA
109141	GCACCCACGC	GTACGCCTCT	AACAACGACC	GGCCGATATC	GTGCAGCCGC	GATCTCACGT
109201	CGCCGTTTTT	AGGTGGGTTA	TCCGGCCTCC	AGCCGGTAGC	AATCTCGTTC	AGGGCGGTCT
109261	GAAAGGATGG	GGCAGAAATT	AACGTGCAAG	CCCATTTTGG	GGGTCGTCCG	TCCCAGGCAC
109321	CGAGCCCGTA	CGTCACAAAA	CACACGTAGC	ATTCGGGGCA	TAGTCCGATT	GAGCGTATAG
109381	AGGCCGCGAG	ATCTAAGCCC	AGCCGAGAAC	CATCCAGCCA	ACGATGGGCA	GGATAAGAAC
109441	CGCGCCCATG	ACAGGCAGCT	TCGTCTTCAG	CCAGGTTAGG	CAAGCGGCC	GAGGCCATCC
109501	CCCAATTGTA	CCGATTGAAT	TGGTTAATTG	GTCATCGCCG	CACGCTCTCC	CGCCCCAATA
109561	TCCTTCAACT	CCGACCCCGA	AGGCGGGGCG	ACGGAGCCCG	TTCGCCTCAA	CGCACCGAGG
109621	CCGTTCGAGT	GACCCGTCAG	GGCAAAAACC	GTTCTAAGAA	GGGTTTTTAA	CCGTTTAGCG
109681	CTCTTTGGAG	TCACGACCAA	AAACTGTAAA	ACCTGTCCGT	GCTCCGTAAA	GTAGGTGCGG
109741	CATATGACCA	TGGAGCTGTA	AACGTTTAGG	TCTCCGGAGA	AAACCAGACG	TGCCTTAAAT
109801	TTCATAAAAT	CGTCCTGGCC	CAGGGACACG	GACGAGTTCC	TCTCAAGATA	CACGTCCGAA
109861	TTTATAGGCA	GGTTTTTTCC	AAACTGGGCA	TCGGCGTCAC	GTGGCTTACA	CAAAAAACAT
109921	TTCAGCGTGG	TGGCCAAACC	GTTGTTGATA	ATTACAAAAC	ACGGGGCAAA	CGGGTAGGCC
109981	AGTCTCTCTA	GTTTGTGGAG	CCAAAACCTA	TACACAAAAC	CGAGATGATA	GACGCAGCCG
110041	TGCTGCAGGC	GCACGGTGCA	CACGGGGACC	GCCCCGCCTT	TAGCGTATAC	GGGAGCCCCG
110101	TCCTGACACC	TCTCCAAGTC	CAGGGAGATT	CCAGAGGGTC	CCAGGTAAGA	GACAACATAA
110161	TCGCACAGCT	CGTCAACTAA	ACGTTTTCCG	GAACCTCATCG	TTATAAAGAT	CCTTTAGGTG
110221	CTGTGCGTGG	CTCCCGTAAA	AACCGCGTCC	GTGCTAACGA	TTTTGTGAAT	GACCTGTTTT
110281	ACGGCGTTTA	CCTTGGCGTC	CAGGACCATG	CAGTGCTCAC	AGTGAGCTGA	CCGCGTCTGA
110341	GCACGATGAC	AGAGGAAAGT	TTTTAAATAC	TGACAGTAGT	TAATGGCGTT	GAGCCTGGAA
110401	TATATGGTGG	GAAACATAAT	TTTCATGTCA	TCGGGCAGCA	GGGACTCGAA	CGCCAATAAA
110461	TCGTCACCGA	ACATCACGTG	AGACAGAGGT	AAAAGATGCT	CACCGCCGGT	ACCGCGTAAC
110521	ACGCGACCAG	CCACCCCTTC	AAATATTTTA	GCCTTAAAAA	GCGGGCCCCC	TAAAGTCGTC
110581	CAACTCAGCT	TAAAAACTCC	TACCCATTAT	GTTCTGTCCT	TGGCAACTGG	AGACCTCAT
110641	GAGACACTGG	CCCTCTCTGC	GGGGACTCGT	AGAACAATCC	TTCTCTCCCG	GTACCCCGGA
110701	CGGAGCTTTT	AACAGCCCGG	TATTAATCCA	CACTCAGGAC	TCTCTACAAC	CCGCCTCATC
110761	GTGCAGGGTG	TGTAGCCTCC	TGTTCACTCT	GGTCCGGACA	TTCCCACCCC	CCGACTCTTT
110821	CTTCGAAGAC	TACGGCTGGT	TGTGCCTCAC	CTGCCTATAC	GCCCCCGGAT	CATGGACGGC
110881	TACCTCATG	GTGGCTGCCG	ACCTTTTGGA	ACTAACGCAC	GTGTACTTCC	CGCAATGCGT
110941	GAAAGATGGG	CCAGTATACA	CCGCCCCAAG	CATCCTCGGA	ATCGACGTCC	AGCTGCACTT
111001	CTTCGCAACC	CGCTGCTTCC	GACCCATCGA	CAGAGAACAA	ATACTCCACA	CATCTCATTT
111061	AAATTTTTTA	CAAACCGAGT	TTATTAGGGG	CATGTTAGAA	GGCACGATTC	CGGGATCGTT
111121	CTGTTTTTAA	ACGTCCTGGC	CGCGCACAGA	AAAGGACGAC	CAACAACCTA	CCGTTGCGTG
111181	TTGTTCCGTT	GGCCGCGGAA	GTCACACCAA	CCGGGATAAC	CGCTACCCG	AGGACCTGGA
111241	AGAGGCGTTT	AACTCCACGA	ACCCGAGGGA	AAAGCCCAGC	CTCCTCGGCG	TCTTTTCGGC
111301	AACGTGGGCA	GAATCCACAG	TTCTTGCTC	CGACACACAA	CAGGCAGATA	CCCATTTACA
111361	ACCCTCCGCC	TTCCCAACCC	CAGAAGATGC	TGACCAATCA	CAGGGCCCCCT	GCCTGATGCA
111421	CCCAACGCTC	AACCTAAAAA	CAAAAAACCA	CACCGCATCC	ATATGCGTTC	TATGCGAGTG

111481	TCTGGCCGCA	CACCCGGACG	CCGGTCCGGT	TCTGAAAGAT	CTGCGTCGCG	ACATTCTGGA
111541	AAACATGGAA	AACAACGTTA	AGCTCGTCAA	TCGCATATCG	TACATCCTAA	ACGATCCGGA
111601	CTCACTGTCA	CACGTGCGCG	ACGAACATCT	GCGCGGCCTA	ATTAAACGGT	GCTCGGCACA
111661	AGAAATCCAC	AAGCATTTTT	TTTGCAGCCC	GGTGTGCGTC	CTGAACACGT	ACTCGCACTG
111721	TCCCGCGGTT	TTATTTAAAT	GCCCACCTCC	CGAAAAGTAT	AAGAAGCTCA	AAGCTCGTCT
111781	GGCAACCGGA	GAGTTCCTAG	ACTGCAACAG	AATATTTGAC	TGCGAGACCT	TACAGACCCT
111841	GGCCGTCTCT	TTTAAGGGGT	CTCAACTGGC	CAAAATCGGC	AAAACCACGT	CGCTCGAGAT
111901	AATCCGTGAA	CTCGGATTTT	AACTGCGTCG	ACACAACATT	CAAATCACCC	ACCCGTTTCA
111961	AACCTCCAAC	CTATACATTT	AATCTTCAGA	AGCGCACCAG	ACAATGCCAA	AACAGCCCCAG
112021	AAGTCGATTG	GCGTCTCGAG	CGCCGTACGC	ACCTAGCGTC	AGGCGACCGG	ACGGGCCCCCA
112081	GTCCACGCGA	CCGGCATCCA	GGCACGGCAG	CTGCAAAAGC	GAAATCATGC	AGTGGAAGAAA
112141	GTTAGTTTCA	GACACGCAGT	TTTTTTCTGC	CCTAACGCGC	CGCCACGAGC	TGGGGGTGGA
112201	CTTTTAAAGA	GAAATGGGGA	CCCCGATATG	CACCTCAAAG	TCCGTTATGT	TGCCGTTAAA
112261	CCTAAAAACC	ATCGCCCCGG	GTCGGTGCGT	CTCTCTCTCA	TCATTGCGAC	ACTCGTCAAAA
112321	CATGGGGTTC	AACTGTTCTG	CGTGACACGC	AACTGACAGG	TCAGCGGTGT	CTCTGGACGC
112381	AAACGCGCTC	GGCGAAGATT	CCGCCAGGAA	AAACAGCGAG	CTGTGTTTCA	TGGCGTTAAC
112441	CTTTTACCAC	CACGCCGAAA	AGGTCGTGCA	GCACAAGGGC	TTTTACCTGT	CTCTGCTCAG
112501	CCACTCCATG	GAAGTCGTTA	GGAAAAGCTT	CACGCAACCC	GGGTTGCTCT	ACGCCCACCT
112561	AGTCCTAAAA	ACCTTTGGCC	ACGATCCTTT	ACCTATTTTT	ACAGTCGATG	CCGATGAGAG
112621	ACTCGCACTC	TGGGCGGTGT	TCCACACTAG	AGACCTACAC	CTGGGGGAAA	CCAGTCTGCG
112681	ACTCATTATG	GACAACCTTC	CAAATTATGA	CATAACGGTG	GACTGCATCA	AGCAAACGTA
112741	CATAATGAAG	TTTACACCCT	CGCGACCGGA	CAACGCAACC	GTGACGGTTC	CTGTCAACAG
112801	CATTTGCGAG	GCCGTGGCCA	CCCTAGACTG	CACCGACGAG	TTTCGAGAAG	AAATTCAAAG
112861	GGGCACGGCC	ATCATAAACT	CCCAGGGGCT	ATTGTAACCT	TTCCCTAGAC	GGAAACAGAT
112921	GTAATTACAC	TATTCAAATG	TTAAGTTTTT	GTTTTGAACA	TATCACCAAT	AAAAACAAT
112981	TATGTTTACA	CAAAATGAAG	AGCGTTTCAA	TTTTACCATA	AACATACCAA	AACCCACGGT
113041	AACTAAAACT	CAATAGCGAT	ATTGCGATGG	GATCCCTAGA	GAGCAATCGA	CGTGCGCCGG
113101	TATTAATTGA	CGGGAACGCA	CGTTAACTGC	TCTCCACGAA	TCGCAAACTC	CGCGTTTTTA
113161	GGACCATTCT	ACGCCGTTAA	TAATTGGCAG	TAAACTGCGA	AAGGGCGTAC	ATCGCAGTAG
113221	TGATTTTACG	ATGTGCACAC	CTTTCCGCGG	TTCACCGCCA	ACAACAAAAG	CGGGTCACCA
113281	GGCACCTAAA	TCGCCCGGGC	CTTTTATGCC	AAAGTACAAA	AGGGACCGGC	GTTCTACGTT
113341	CACGAACAAC	ACTGTTGATC	GCAAATACAA	TAGTGTTGTT	GGTGACCGCA	AACTGCCAAA
113401	TCCTAAAATG	CATGGACATA	GTCCTCAAAA	TCTAGAACGA	ACTTCCAAAA	ACATGCGGCA
113461	ATAAACCAAC	CCTTGACAGT	AATGCACATC	GTAACGAGAC	GGAATCCAAA	GAACACGCCC
113521	GATCTTAAGA	CCACGGGCAC	GTGTCTTTGG	GTTCCGGGGG	CGTTACGATG	GAAGTTCATC
113581	ACCAAGTATC	CCATCAACCT	GTTACAAAAG	CGTAAAGATA	ATCGGTTTAC	AATAAACAAAT
113641	AAAAAATAGT	GGACGAAGTT	ACATTCCGAC	ATACAAGCGG	ACCCAAAAAA	ACACCCCAAC
113701	GCAAAACAAC	TGGAGACTGT	GTCTTTGGGT	CCCTTCCACG	TCGCAATGAG	ATTTCTGGTG
113761	ACCGTGTGCT	GCTTAACGCG	TTACAGTTGT	GTTTATGCG	TACGTAAACA	CACCAATCTA
113821	GAATGCTAAC	ACGTGCACCG	CGCTTAGGCG	CGTTTGTGCA	ACTAACTGCT	GTCAGGTTAC
113881	TACTCTTATT	TTTAACAAGT	AGCATACCAA	CTGCAATTAA	TTGCACTAAA	CCCAGTTTAG
113941	TCTTTTATAG	CGAGTCGGCC	CATAGTCTTC	ATGGAGAACC	ATCATTTCAT	TCCGTCAACG
114001	GCGTCGGTGT	TACCGTTGTT	ATTGCGGGCA	AAACAGGCTC	GGGGGAAAAA	AGAGCAACAT
114061	ATCGTCAGTT	TAAAACAAC	CAAGTTCTAA	AATCTCACCC	ACTCGTAAAC	AAAAAACATC
114121	CGAGTTAAGG	TGTATGTACC	GAACATAACA	ACACAAGTTT	TTTAAACAC	AGCTGCGGTA
114181	AGTAACCCCC	ATTGCCACGC	GTGTGCGGTG	CTAAGTGTTT	TTAAATTTAC	ATTGTGCGTT
114241	TTACACACCG	AGCAGTAATC	TCAGGAGGGC	GGTTAACGAG	CGATATACAT	ATTCCCTAAA
114301	CACGGGAACG	CGCGCTGACC	GCCCTCCCCA	AATCACAACA	CGGGACTACA	AAGCCTAGTG
114361	TTAATATAAT	CAAATTAAAA	AACCACAGAA	ACCTTTAGTC	GTGCGCAAA	ACTAGCAAAAG
114421	GTACCTAGAG	CTTTCCCCTA	TACTTCAAAA	AACAGCGGTG	GGTTATTTGA	CCACACGTTA
114481	AGTAAACACC	CGTAAGAATT	ATTCCCGTTT	ATCAAAATGG	AAAAATTAAG	GCTTTGCGTT
114541	AAAATCTGCT	AACGCAAAGG	GCACTTAATT	TTTCCAGTTT	GGACTCGGAA	CTTTACCGCT
114601	AACGTTAAAA	TTTAATAATG	CAAAAGGCAC	ATAATTTTTT	TTGTGTAACC	TCAAAACTCT
114661	ACAGCTAAAG	TTAATAACGG	CGACTTTGGG	CAGCGAATCA	GTGTCTGTCT	CAAACCTGCT
114721	TAAATTTAAA	CACAACAGGG	AGCGCGACTA	AACACGGACT	AACTGCTCAT	CGGGCCCTGT
114781	TGAAGAAGTT	GGTGTGCAAA	TGCATTAGGG	AATCTAAAAA	CGAAATCCTG	CTCGGTAACA
114841	CGGAAATTGT	CTTTAGTTCC	CTACAGCATC	ACAACAATTG	TAAACCATAA	ACGTACGCGC

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114901	GTGGGGTTGG	TGTGGTGTGA	AATGCTTGGT	GCAACTGTTA	CATGGGCGGA	TTGTAAATGT
114961	GGTGCTTGGT	GCAACGGTGG	TGGGGTGCAA	GTCCCCCGGT	GGGGTGCAAG	TCCCCCGGTG
115021	GGGTGCAGGT	CCCCCGGTGG	GGTGCAGGTC	CCCCCGGTGGG	GTGCAAGTCC	CCCCCGTGGG
115081	TGCAAGTCCC	CCGGTGGGGT	GCAAGTCCCC	CGGTGGGGTG	CAAGTCCCCC	GGTGGGGTGC
115141	AAGTCCCCCG	GTGGGGTGCA	AGTCCCCCGG	TGGGGTGCAA	GTCCCCCGGT	GGGGTGCAAG
115201	TCCCCCGGTG	GGGTGCACGT	CCCCCGGTGG	GGTGCAAGTC	CCCCCGTGGG	GTGCAAGTCC
115261	CCCGGTGGGG	TGCAAGTCCC	CCGGTGGGGT	GCAAGTCCCC	CGGTGGGGTT	CAAGTCCCCC
115321	GGTGGGGTGC	AAGTCCCCCG	GTGGGGTGCA	AGTCCCCCGG	TGGGGTGCA	TGCCCCCGGT
115381	GGGAGCGGCT	CGGCTCCGGG	GTGGCTCCGG	GTGGGGCGCG	CTCGGCTCCG	GGGTGGCTCC
115441	GGGTGGGGGC	GGCTCGGCTC	CGGGGTGGCT	CCGGGTGGGG	GCGGCTCGGC	TCCGGGGTGG
115501	CTCCGGGTGG	GGGCGGCTCG	GCTCCGGGGT	GGCTCCGGGT	GGGGGCGGCT	CGGCTCCGGG
115561	GTGGCTCCGG	GTGGGGCGCG	CTCGGCTCCG	GGGTGGCTCC	GGGTGGGGGC	GGCTCGGCTC
115621	CGGGGTGGCT	CCGGGTGGGG	GCGGCTCGGC	TCCGGGGTGG	CTCCGGGTGG	GGGCGGCTCG
115681	GCTCCGGGGT	GGCTCCGGGT	GGGGGCGGCT	CGGCTCCGGG	GTGGCTCCGG	GTGGGGGCGG
115741	CTCGGCTCCG	GGGTGGCTCC	GGGTGGGGGC	GGCTCGGCTC	CGGGGTGGCT	CCGGGTGGGG
115801	GCGGCTCGGC	TCCGGGGTGG	CTCCGGGTGG	GGGCGGCTCG	GCTCCGGGGT	GGCTCCGGGT
115861	GGGGGCGGCT	CGGCTCCGGG	GTGGCTCCGG	GTGGGGCGCG	CTCGGCTCCG	GGGTGGCTCC
115921	GGGTGGGGGC	GGCTCGGCTC	CGGGGTGGCT	CCGGGTGGGG	GCGGCTCGGC	TCCGGGGTGG
115981	CTCCGGGTGG	GGGCGGCTCG	GCTCCGGGGT	GGCTCCGGGT	GGGGGCGGCT	CGGCTCCGGG
116041	GTGGCTCCGG	GTGGGGCGCG	CTCGGCTCCG	GGGTGGCTCC	GGGTGGGGGC	GGCTCGGCTC
116101	CGGGGTGGCT	CCGGGTGGGG	GCGGCTCGGC	TCCGGGGTGG	CTCCGGGTGG	GGGCGGCTCG
116161	GCTCCGGGGT	GGCTCCGGGT	GGGGGCGGCT	CGGCTCCGGG	GTGGCTCCGG	GTGGGGGCGG
116221	CTCGGCTCCG	GGGTGGCTCC	GGGTGGGGGC	GGCTCGGCTC	CGGGGTGGCT	CCGGGTGGGG
116281	GCGGCTCGGC	TCCGGGGTGG	CTCCGGGTGG	GGGCGGCTCG	GCTCCGGGGT	GGCTCCGGGT
116341	GGGGGCGGCT	CGGCTCCGGG	GTGGCTCCGG	GTGGGGCGCG	CTCGGCTCCG	GGGTGGCTCC
116401	GGGTGGGGGC	GGCCTAAAT	CCTTACCGGT	AAATTTAGCA	GTAAATCCAA	CGAGTAAAT
116461	CCGCAAGCTA	GCCGCACAGA	GCTGCGACTG	CCTGCCAAGG	CTCCTGGCGC	CTCTTTTATA
116521	CCGCTAAATG	CCCTCCCCAA	ATGGTTACTA	TGGTTTAGTT	ATGCAACCAT	AATACCAATA
116581	AATGAATCAC	CAAGAATACG	GCCAACGCGC	CAGCGTCCAA	CCGCCCTCCA	ACCGTGCAGC
116641	GCCGTATATT	CGAATTGCCT	TCCGCACGAA	TACGGGGCGC	AGCATGGTCG	CGGGAGGCTG
116701	GCATTGCGCA	ATATTCCCAG	CATTCCAACG	GGCACGCAAG	GGTCCAGGCT	ACCGGGGATT
116761	CCCCAAACAT	CATCCAGCGC	ATAGAGGTTG	TGGCTGAGCT	GCCTATCTGG	GGATTCCCCA
116821	AACCCGAGAC	CCACTGAGAT	GCTATTCACT	GGGACTGCAG	GCTGGGTCTT	CCCAGGACAG
116881	GTGCCTGCGT	GTGTGGTAAG	GGAGTTCCCC	TAATTAATAA	ATTATATTAA	TAAATAAACC
116941	AGGCTAAGCT	GTAATTAATG	AGGCACAGGC	TGCCGACCTA	GGTACCTGGG	GAACCCCTTA
117001	ACGGGATGTA	ATTAATTCAA	ACCTATATAA	ATTCCACCCT	GTTAGGGGGA	TCCCCACGTG
117061	TACCTGAATT	AATGCGGGAT	ACAGGCTGTA	GGTTTGGTAA	TGGGTACCAG	GCTGGAGTGA
117121	CTCATACCAT	GGGATTAAAT	ACCAGGGATA	CAAGCTGTAG	GTTTCGGTAAC	GGGTACCAGG
117181	CTGGAGTGAC	TCATACCATG	GGATTAAATA	CCAGGGATAC	AGGCTGTAGG	TTCGGTAATG
117241	GGTACCAGGC	TGGAGTGACT	CATACCATGG	GATTAAATAC	CAGGGATACA	GGCTGTAGGT
117301	TCGGTAATGG	GTACCAGGCT	GGAGTGACTC	ATACCATGGG	ATTAAATACC	AGGGATACAG
117361	GCTGTAGGTT	CGGTAATTTA	AACCTATATA	AATTTTACCC	TGTTGGGGGA	ATCCCCGTTT
117421	GTACCTGAAT	TAATATAGGA	ATACAGGCTG	TAGAATCGGT	AATGGAAACC	AGGCTGTAGG
117481	ACTAGGCTGG	AGTGACTCAT	ACCATGGAAT	TCAATTAACA	AGGAAATAT	AATAGAATAT
117541	ATATATATAT	AGGGAACCTG	TAAACAAAAC	CCAACTCGCG	GATTGGCTGC	CTGTTTGGGC
117601	CAACCAGCAG	CGAGAATTCC	CTGATTGACA	GGCGGACTGG	CCAATGGTTT	GCGAGCATTT
117661	TTGATTGACA	GAACGGCCGG	CCAATCGCAA	CCGAGAACTC	GGCAGCGAAG	CAAAAGACAG
117721	ACGGCCGCGG	CGACCAATGG	CCGCCGCGGG	TTAGTTTGAT	TGACGGCTTG	GCCGGCCAAT
117781	GGGAAACGTC	CCTGCCGGCG	GCCCCTAATC	CCCCTGGCTT	AAGGGATTAC	GGCCGTAAAC
117841	ATCGCTGGCG	CGGTGCCACC	GCCGCCGAAC	CCCCGCGCCC	GGTTCGGCGT	GACCTCGCCG
117901	CGACCCCGCC	GCCCTCATGC	GCACCATGGG	TGCGATGGGC	ACCGGCGGCA	ACATTATTCG
117961	CTGACCGCCG	GACGACCCCG	GCGCCAAAGA	GCGGCCGATG	GGTACGCGCG	GCGACCATCG
118021	AAACATTTTT	CAATAACCCC	AGTCTGACCC	CGCGGCCCAA	ATCTGGGGCC	CATGGTGGCG
118081	CGCGGCCCCC	ACCGTCCCCA	TTTCCACGCG	TTTGCTCATG	GGCACCAGCG	GTGCCCCGTG
118141	TGATATGTTT	CAGTGACCC	CGATGACCC	GGGCGACGTT	TTGCGCACCA	TGCTCCCGAA
118201	TTCCCCCATG	AGTACATGTT	TCGGTGACCC	GCCGGCGACC	CCGGGCGGCA	AAAGCCGCCC
118261	CATGGGCACC	GGACGGCACC	ATGTGCACAT	CTTTCAGTGA	CCCGCGGGCG	ACCCCGGCGG

118321	CGGCACCGCG	GGCGCTGGGG	GTGATATGTT	CTGATGACCG	CCGCCTTACG	TTTGCAGCGG
118381	TGTCAGCGGA	ACATATTCTC	CCCGGCTCCC	GAGCCCGGGG	CCCGCGGACC	CGAGTTGGGT
118441	TATTAAAGTT	ACGTTAGGGA	TTAATTTTAA	TCCACTACCC	CCCTTAATGT	GCGGCTCGGC
118501	CTGCGGCGAC	GCTGACGCCG	GCGGGTTCGA	CCCATTGACC	GCATCCTCGG	TGGCCGGGCA
118561	GAGGCCGGCC	GGCCAAAGGT	GCGATCGCGG	GGTCTGACCC	ACCCGTGGGA	GAAACCCCTC
118621	CGTTCCGTTG	TCTTCTGCCT	GCGGTCCCTG	GCCCCCTGGC	CCACCGGGGA	TGTCTTGTGA
118681	GCACTTTCCC	CGGTGAGAAT	AGACAGGAGA	GTGCCCAGAA	GACATCGACG	GGGTGCGCGG
118741	TCGGTGAACG	GGAGGCGCGC	CGCTTGCGTG	CGTGAGGTGT	CGCCGGTGGG	CCGTATGACC
118801	CCCGCAACGC	GAGGGGCCGC	TGGCGCGGAA	ATTTCCCCAG	AACCGGCCGC	GCCGCTTGGC
118861	GGCGCGGCTT	CCTCCCCCAA	CGACCCCTAA	TTTGGTTTTT	AAGTTGTTAT	TTAAAGCTCC
118921	GCTGTGTTTG	TGGCACGCGC	TTAAGCGTTG	CCACCTGTTG	CGGAGATCCC	CCGTGCGCAT
118981	CGCCGCCCTC	CTCGCCTCGG	TGCACGCTGC	CGAGCGTGCC	ACCTGTTTGG	TTGTGTCTATC
119041	CCATGTGCGC	ATCCCCCAT	CTTATTTCCC	CGACCAGGCA	ACATAAACGT	CACGTCGTTT
119101	GTAAGAAATA	ATTGCTTTTT	TTATTTTTCGT	TAACCCCGCA	ACCGCGATAC	AGTCTCTCGC
119161	CTTTCACCCG	TTGTTTTGTT	ACCAGCCCCT	CTTGGCGGCC	GGCACC GGCG	TTAACCGGGT
119221	GCGTTGGCGG	CCACGACCTT	GGCCACGTCG	CGCCTGGAGA	CGGCCTGCAG	CAGGTCACTT
119281	AAAACCATGT	AGTTTGAGGG	GCTGACCAGG	GCCGCCTTTT	CCATTTGCGA	GAGCCACCGA
119341	AAAAAGGTGG	GTGACTGGTT	GCGCCTGCGA	CCAAGCTGGT	CGCCCGTTAG	AAAAACTAAA
119401	TTTTTCACGT	CCCTTTCTGA	GAGCTGGCGG	TCAATGGAGA	GCATCAGGTG	TTTGTATGGA
119461	CTAAAGTAAC	TGGGGCTGGA	CCGGAGCCTC	TCGATAATAA	ATCCCACGTC	TAGTAAAAAA
119521	AAGGTGGAGA	CCAGGTCCAA	TCGCCCAACG	AGAAACACGG	CCTCCAGCAA	CACCGGGAGG
119581	GGAATACCGG	GTTCTCCAGT	TGAGGGGCAC	AGCCCGTTTG	CAAACCCCTC	GGGCGTATCG
119641	TCGGACGCCG	GCCGGTCAAA	AAGCCACAGC	ACGGCCTCTC	TGTCGTCTGC	CTCTAGGTGG
119701	CGACCGAAGT	CCACTAACCG	CTTATGCGGG	AACATGGGTC	CGGTGCGGCG	CTCGCCGACA
119761	CCCTCTTCTA	CCTACACTAG	AGCGTTTGAT	AAATCACGAC	GAACCGCGGG	CGTTACTTAA
119821	AATGTGGTCT	TCGTCAAATT	CCCGCAAGGT	GGTGGCAACC	GACTCGGCGG	CGCCCGCGAG
119881	GGTTGCGGCC	GAAACGCCGA	CGGCCTCCGC	CAATCTGGCC	GGAGCGGGCG	CAGCGGCACC
119941	GAGGCTAAAC	AGCGCGCAGG	CGGCCGCCAC	CAGGGACGGG	GGCAGACCGC	CGGTGGCCGG
120001	GTTTACGATG	GCTTTGTGAA	CAGACTCCAC	CACCTGGCTG	TGAAGGGCAA	AGAGCTGCTC
120061	TTTTGTAAAG	CCGCTTTTAA	ACAGGGTGGG	CCCGATGGCG	TCCGTTGGTA	AGACGGCCTC
120121	TAGTTTCCAC	CTTAAAGCCT	TAAGGATTGA	TTTTTCCTGT	TTCAGTAAAT	CCGCCACGGA
120181	AAAATCCTCC	GCCGCGCAAA	AGCATAAAAA	GGCGGCCTTG	AAAGGATTCA	GGTCTCTAAT
120241	CTTGCCGGCT	AAAAATAGAC	AGGCGGCCCC	CAGCCTCTGA	AACCGCCGAC	GGGGGATGCT
120301	GCGACACTCT	AGATAGCGAT	CCAGAATACT	AACGGCCAGT	GGGAAAACGC	TAGCATCCGC
120361	CTGGTGGGCC	CTGGCAACAG	ACCGCATCCA	CGTCCCCAAG	ATAACTCGCA	TGCCCCACGT
120421	CACCTCCGTT	TGTATCGTCC	CATAGCAGGC	CGTGGATGTT	ACAAAGGACG	CTTCGTGGGC
120481	CAGGAGGTTA	GAGAGGGCCC	GGTCTTGTTA	CAGGACGGGG	TCAATGGTCC	CGGTGGGCAC
120541	TGGGCCAACA	GAAGCCATTG	TCAAGGCAAG	GTAAAGAAAG	CGATATAAGT	CCTACCTGCG
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120721	GGACTAGATG	CAACGCGGAA	GCGGCTGTTT	ATTTAAGGTG	AGTCACAGGC	GGGGCAATTT
120781	GCCAACCAGT	GACGCGTTTC	CTGCAGTCTG	GGTTTTTACC	TGGCAAGCGC	CACTGGTTCT
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121021	TGACGGGACT	GCCAGGTGCG	TCCGGCAAAA	CGCGGACGCT	GGATGCTCTA	ATGCCGGGAA
121081	ACTGTAGGCA	CCTTGCCAGG	CGATACAGTG	AATAGGGGTC	TGAGTTACAA	AAGATGACTC
121141	CGTGACACCA	CGGGCCCGGC	CGAGGGGGGT	CGATCCTGGG	TCCCAATACC	TGACGATAAA
121201	ATCTTCTCTG	TGGGCGGTAT	TTTCTGGGTA	CCTCTTCCTC	TGATGAGGGA	TACGGTATGG
121261	GAATGGGGTA	TCGCGGCGGG	CGCGGGGGGC	GCGGGGGGCG	CGCCGGTTGC	TGCTGCCGCT
121321	CCTCCCCCTC	TTGCTCCTGT	TGTCTTTCAG	GAGCTTCGTC	CTCGTGGGAG	TTGTGACTCG
121381	TGCATCTGAG	ACGTAAACAA	GGAATCCTC	CGGCACGCGT	GGGTGGACCC	CAACCCCTTA
121441	CGGTGTACGA	GTGGGATGTA	TGGCCAGGAA	ACGGGGGTCC	TGGCCGAGAC	GGCCCGGTGG
121501	GGTGGTCTGT	GTGGCTCACC	GCCTGTGTGC	TGTTAGGGGA	TGGGGCCTGT	TGAGGCGTTG
121561	GCGATTGGGG	TGGTCCCGGT	GAGTCGGGCG	GGGATTGCGG	AGGGGGCGAG	GACGTCGGCG
121621	TTGGCGATTG	TGGCGGTCCA	GGTGAGTCGG	GAGGGGGCAA	GGACGTCGGA	GGCTCAGGCG
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121741	GATTATGCGA	CGACGGCGGT	GCAGGTGAGT	GGGGGCCGGG	GTGTGACGGG	GGTGATGGTG
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121861	CTGAAGCTAA	CACCGGCGCA	GACCCGCTTG	GCGACGGCGT	TGGAGATCCC	AACACGATGA
121921	CGGACTCAGA	CCCTGACCCT	GATGGCGACC	CTTGGGATGA	CCCGTGCCCTG	GATTCGGGTGG
121981	CCGTTTCGGT	GTCCGATCCG	TGCCCTTGTT	CTGCAACCGC	CCGCACTTGC	GGCCCTGATG
122041	TTGTTGGAGC	GGGGGGCGCG	GGTGCGGCGG	CTGGCGCGGC	CGTGCCACGC	CCCCTGCCGC
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122161	GCAACCCGTG	TCCCGATACT	ATGCCGGAAC	GATGTTGCCG	GCTGCCCCAC	ATGTTCCGCG
122221	TTTACTGCGT	GTAAAAACG	GCAGATTAAG	TAGATTGCCC	TTTCCACGAC	CCGCAGCCAC
122281	CTGAGCGCGC	GCACGCAGAA	CCTATGTTTA	GTGCGATTGC	GATTATGTCC	GCTAGGTGGC
122341	GGGCTATTTT	GATCCTGCCG	CGCACGCATT	GACCGTTAGG	TGGCGCAGAG	CGCCGTTAAC
122401	CGCAGCGAGC	ACCGCACGGC	GGTTTTAGTT	TCGTTACTAT	AACGTTTAGT	TAAAATCCGT
122461	GGCCGATATT	GGGTGGCAGT	GTGTAAGTCA	CGCGCGCCGC	GGCCCCCTTA	CCTTGTCGCC
122521	GGCGAACCGC	GGAAAGTCTG	TTCCGCCGCG	CAGCGCGCGG	CCGCTGCCGA	AACCAACGCC
122581	GCTTGAATTT	CGCGGGCGGC	TGGGCTTGGT	GTCAGTGATT	CACACAGATG	CGATGTGATT
122641	GGGCGTGGTG	GTCAACGTCA	CTTGTTAAAC	CGTAAATCTG	GAAATCTTGT	TCCGCCGCGC
122701	CGGTGAGCCG	CGGCCCTAAG	CCTTATAGTG	CTGTGCAGCG	ATCCTGAAAC	TCGAAAGCTG
122761	GCCATAATAA	CCCCAGTATG	GCCTTGCGTG	TCGGTGGCAA	CCTTTTGA	AAAGATTTAT
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122881	ACATTAACGC	TGCTGCTGGC	GACGCTTGCC	ACGGTTCGGT	GCGCTCTTCA	AACGCATAT
122941	GCGGCGGTCC	CCGTGCACTC	TACCGCGTCC	CTGGGGTGCG	TGTTAACGAC	ACCCACGAC
123001	GTTCTTATCG	TTACCTGGCA	AAAACAGGAA	TCGCCTAGTC	CCGTTAACGT	GGCCACATAT
123061	AGTTCCGAAG	CGGGCACGGT	GGTTCAGCCC	CCGTTCGCCG	GTAGGGTTGA	CATTCCCGAA
123121	CACAAGTTGA	CCAGAACGAC	CCTGAAGTTT	TTTAATGCCA	CCCTGGAGGA	CGAGGGGTGC
123181	TACCTGTGTA	TCTTTAACGC	GTTTGGAGTG	GGAAAGCTGT	CGGGAACCGC	CTGCTTGACG
123241	GTTTACGTCC	CCCTGTCCAT	GTCCGTACG	TTTTACCCCC	CGATTAACCC	CAGCGAGCTC
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123361	CTGTGACGCG	AACCTGAAGT	GTTTCCCGCG	CCCAACGGAA	CAACCTGGT	TGTGGGTGCG
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123481	ATAGGCGGTT	TGGCCGCGGC	GCGGCCCTG	GACCCCGTGT	TTTCGGATCC	CCTGGAAGGG
123541	ACGAGCCACT	ACGTGGTGGG	TGTGGTGGCA	GCGGCCGCCG	TTTTAGGCAT	TTTTTTAACG
123601	GGTGTTTTTT	TGTATAGGTC	TATGTGAGCG	CGCGTGTCCC	CCGTGTCTAG	TGTTTTGTTC
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123721	GTGCGACGCT	TCCTGTGTAA	CTGCATACAC	CGGGGTGTCG	CCAGGAAACC	GCGTCTCCCC
123781	TTTATGTCCG	CTCGCCCTCC	CAGACGAAA	GTGAGAATGG	TTCTGGGGC	GTTTTGGCGT
123841	TGAGAGTGC	GGGCGATGTT	GCCGTAGCGG	CGTCTGCAAA	GGCTCACCCG	CTTCTGTTTT
123901	TTTTCTTTTT	GTCAGACAAC	AACATGGACG	CCTTGAACAA	TAACCTTAAC	CTGCTGATGG
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124201	ACTCCCTGTT	TCTGTGCGCG	TCGTTTTTGT	TCAGCTGGCT	GCTGTACGTC	GCGCCACAGA
124261	TGCTCACGTC	CGCGACGTGC	AAGGTGGAAA	TCTTTTCTT	TTACCTGTAC	ACGTACTTTG
124321	GCGTGATCAT	TGTGGTGTGT	ATCAGCCTTA	TCAGGTGCCT	GTTAGTTGTG	TTTTCCCGCC
124381	GCCCCGTTGG	CAAGCACGGG	GCCTCCGCGT	TTCTCTGCGT	GTGTGTGTCT	TTAATCGTGG
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124501	GCGAGTGGAT	ATGCTACGAA	GATGCCGGGG	AAGATAACCGT	CAACTGGAAG	CTGAGAATCA
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124681	TAATTGTGAC	GGTGGTGGTG	CTGTTTTTAA	TTTTTTGCCT	GCCCTATCAC	CTGTGCAACT
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124861	TCGTGTATTC	TGGTCTTGGG	TCTCTGTTTA	GGAGAAGGGT	TAGGGATACC	TGGTCCGTGT
124921	TTAGGTGTTT	TTCCACTTCA	GGTAGTTTTA	GAGACACTCA	CGCGACACTT	GGTTGGATTG
124981	TTTGTGTACA	TTTATTTTCA	TTTTGTGTAC	ATTTATTTTC	ATTAAAGCGA	TCTGACCTGC
125041	AGACCTTACC	TGACGTTTAC	TGTCTGTTTC	TTATGCACCA	GAGGAACAGG	GACTGGAAGG
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125281	TCGGGTCCGC	GTTGCCCGTG	TGGTAGTTCA	GGGCGATGCG	CTGCTGCTGG	TTGAGATGGT
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126241	ACACCAGCCA	CTGCAGGGGC	TTTTCTCTCT	ACGGGATGCG	AATCTGTAGG	CCTCGGTTAC
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128281	AGCTCCTGCG	GGCCATGACG	GCGTATCTGG	ACACCGTATA	GCTGTTGCCG	TTGGAGGCCT
128341	GCACGGCCAG	CGGTAATATG	TTCCGTTGTA	ACGGGAGCAT	CACCGCGGCG	CATATTGGGT
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128461	CGCGCACATC	ATATTCTACT	AGTGTGCCAA	GTGTCAGGGC	GGCCAAGGTG	CCCGGGGGTA
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 131941 CTAGGGTTAT AGTTATAGTG TAAAAATAA GGCCTACGAT AAAATGTCAA CATATATTTT

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132001 TTAATTTGTG TTTCATTAAC AGCCATGGTG TATTAGAGTA AAGGTTAACG CTTCAATACA  
 132061 ATATACAAGT AAAAGTAAAC CACACGAATT TATAACATAT TTACAAAAGC GAACCAAGCT  
 132121 GGCACATGTA ACTTCACGCT ACCCTAAACG TACAATACTG GGACTAGAAC CCAGAGGTAG  
 132181 TTAGAATATA CGGTAGTTAC AGAACTTTGC AGTTCCCTTA GGCCAGCAGG GCTCTGCGGT  
 132241 TAATTAAACA AAGTTTAAAG ATAAC TGAAG CTTTAGGAAG TGCGTATGGG TGCAATGTGT  
 132301 TCCAAATAGG GCAAGGGTTA CATAAACTGT TGCCTAGCGG CCGGGCCCCG AGGCGCCCCG  
 132361 CCGGCGCCGC CGCCGGGCCG CGGCCGCCAT CTGCGCCCCG GGGCGAGGGT CCCC CGCGCG  
 132421 CCCCCCGGGG CCGCGCGCCG GCGCGCCGCG GCCTCCCCCG GCTCCCGGCC CTCCGCCCCG  
 132481 CCGGCCCGGG CCGCGCGCCG CCGCCCCCGG CCGCCCCCGG CCGCGCGCCG GGCGCGTTTC  
 132541 GGTTCGCGGG GCCGGGGGTC CCGCGGGGGG CCGGGGCCGC CCCC GGGTGC CGCTCCGCGG  
 132601 GGCCCGGCCG ACTCCCGGGA GCGCCCCGGT CCGGCCGAGC CCGCGCGCGC CGCGCCCCGA  
 132661 CGCCCCCGGG GCCCGGGGCC CCACAAGCCG CGGCGCGCGG CTCCCGATGC CGGGCGGCCG  
 132721 CCGCCCCGCA TGGCGGTCTT CCGCCGGCCT CCCCTCCCCC ACGCCGCCCC GAAAGGTGGT  
 132781 CTCCGCGCCG CCGGGAGGGG GGCCGGGGCC CGGGGCGCCT CGGCGGGGCC CGGCGCGGGG  
 132841 CGCGACCGAG GGCCCCGGGA GAACGGGGGA TCGGGAAAAC GCGAGGGGAG CCGGGGACAG  
 132901 GGGACGGCGT GTGCGTGCTT GTGAGACACC GGTACGGCT GCCTGCTGC CTCTGGCCT  
 132961 GCTTGCTGAG GGGACAGTAG GCCTGCTTGC TCGCTGGCCT GCTTGCTGAG GGGACAGTAG  
 133021 GCCTGCTTGC TGAGGGGACA GTAGGCTTGC TTGCTCGCTG GCCTGCTTGC TGAGGGGACA  
 133081 GTAGGGCTGC TGGCTTGCTA GTAGGGCTGC TCGCTGGCCT GCTTGCTTGC TCGCTGGCCT  
 133141 GCTTGCTTGC TCGCTGGCCT GCTTGCTTGC TCGCTGGCCT GCTTGCTTGC TCGCTGGCCT  
 133201 GCTTGCTTGC TCGCTGGCCT GCTTGCTTGC TCGCTGGCCT GCTTGCTTGC TCGCTGGCCT  
 133261 GCTTGCTGAG GGGACAGTAG GGCTGCTTGC TTGCTAAGGG GACGGTACGC CTGCCTGATG  
 133321 GCTTGATAGT AGGGCTGCTG GGCTGCTAGT AGGGCTGCTG GGCTGCTAGT AGGGCTGCTG  
 133381 GGCTGCTAGT AGGGCTGCTG GGCTGCTAGT AGGGCTCCTG GGCTGCTAGT AGGGCTGCTG  
 133441 GGCTGCTAGT AGGGCTCCTG GGCTGCTAGT AGGGCTGCTG GGCTGCTAGT AGGGCTGCTG  
 133501 GGCTGCTAGT AGGGCTGCTG GGCTGCTAGT AGGGCTGCTG GCTGGCTTGC TTGCTTGCTT  
 133561 GCTAGTGGGG CCGCTTGCCG GCTACTAGGG CTGCTGTGCA GCTGGGAGAA CAGAGTAGGG  
 133621 CTGCCGGCCA GCTGCGTGCG AGGGCGTCCG AGGGCCAGAC GAGGACACGG GACCCGGGCC  
 133681 TCTCCCCCGC CCGGACCGCC GGGCACCCGG CCCAGATCT

## SEQ ID NO 2

CDS nucleotides 1353-2624

## SEQ ID NO 3

MFVLVLFMLLQPVSVLLPAKLTSVPTWCPHPGDTYLLTCRGSTARDQRSTQWFRNNTLMRGSNFYGRVSVTPNATI  
 SDRYACQTKTTTRSNNIDFRVSSSRLTLQERCSSYGYTYANNTRVLRCSGGNVTLRNVVHFLNGTAVINGTTTNIHTFV  
 LTEKTGGTYFCSAFIGNKXFYSQTINVFSTFTFKPTNDIPNESHFNKTGQIQQTASVQHPENYVVFVSVFVSIGVLTGI  
 AISLIMCWLFTIRCENSESSTNSYASQTSYIQPSHNQSRNTNECSRHTYRNAHQEESIEELPNQHTSETDSCCQLVLE  
 VKNVAYDGPQENTINEVMEQYDDVVVKNIEQTSYEDNVEHMDYSDTINPNFNYSGLILEEVDEVFYNELNQYHGLILE  
 NLDHNEYNHLNELNMIEQYDWLE

## SEQ ID NO 4

CDS

complement (2692..3258)

/note="dihydrofolate reductase; ORF 2; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 2"

## SEQ ID NO 5

MDITVNCIVAVDEQLGIGKNGTMPWPYLRNEMMYFQKMTSTPSVVGKENVVIMGKRTWFSIPEKKRPLVNRINIILSREL  
 REPPHGAHFLARTLDDAFNFYRQYKLKEQLNTVWVIGGKSVYESVLNYKCPLKLYITRIMESFDCDVFFPSINFTEYTML  
 SEIPGKDTNFEENGIKYKFQVYEKNFNK

## SEQ ID NO 6

CDS

3676..5613

/note="complement binding protein; ORF 4; similar to  
 Kaposi's sarcoma-associated herpesvirus ORF 4"

## SEQ ID NO 7

/translation="MTFKLFPLFLLHAIMVHCDENCKPPHFTEYRVKSNTEKDLYSV  
 GETAELICRPGYVTNTKIITTECLONGTWSTPNFPDCDRKRCPTPADLLNGAVHIHGGD  
 NALKFGSNISYECNEGYDLIGSNVRFILQDTENVNWDSENEPVEIQKCIKPPAVEHG

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DYLPNQDVYNYGDAITFKCSLSYTLVGSTTLVCTSNKKWSNSFPTCLMLVCESQIDN  
 GYIDIGLSRRYNHGQSITVKCSGDYINIVGPETLTCTNTTWPPLPKCVLVTNNPSTPM  
 PETPMPETPTPDYQKINLSTAKTATTNPAFVTTVVSPEKDDVTCVKPHFERFMVKAEN  
 DKEKYSVGASVELICRPGFTKMQSTVSVECLSNGTWTAPNAKCHRRKCPPTQELLNGE  
 YIVTSGEDAFKYGTNITYKCNQYQLLGSMVRI CMLKDDLKTVDWEPKAPICDIEKCK  
 PPPQITNGKYHPVKDFYQYLDTVTFSCNRDFS LGVDEM TTCISNTWNKFPFPRCEQITC  
 SAPNIAHGKLLTGSSSVYKYGQSVTIGCETGFTLIGSEISTCKDSSWDPLPTCVPÄV  
 SMPSDTPKPKETKKPNTPTPEAPKPNTPNVGTHTPFKPPPQNPPPIAPPMSKWKRHVVLV  
 LFASVASLLFVLAALYCCFLK"

SEQ ID NO 8 CDS 6045..9443  
 /note="ssDNA binding protein; ORF 6; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 6"  
 SEQ ID NO 9 /translation="MASKGNAGQPLEDNQGSRAPIGACGYVYAYSKQDFPFAEASILG  
 NRPSGSGVFSLPILYGLTVEHEFPLTVKAAAYKKVDTTTAVKVTCTFHREVIVFHNASL  
 FRPVFDGTGLNELCEEARALFGYTQFIEPGPPHSIWNPLECPQLPDKDEMFLGVVTE  
 GFKERLWRGCLVPAVFQTTQVQIAGRQAFKVPPLYDEDLFAPHGHRMPRFYHKDVSAYL  
 YDSLFTSIAQALRLKDVTAVIDHATEKQFMQDHYKIAKIVQAKQFSTTLPKTTDGSSHM  
 IVDSVVAELALS YGCMFLECPQDACELLNYDSWPIFDGCDSPPEARVNALERWSAEQAV  
 HVAGQLFAANSVLYLTKVQKQAPRGQKGDVNVNSFFLQHGGLGFLNEATIKENGSEAF  
 KGVPSNALDGSSFTPYHLAYAASFSPHLLAKLCYMQFLQHHKSSTNQAFNMVHYVGT  
 AANSEMCTLCHGNTPATCLNTLFYRLKDRFPAVTTPORRDPYVVTGTAGTFNDLEILG  
 NFASFRDREEDGNPADEHPKYTYWQLCQTVTEKLSAIGITEDHDNHVNLTNIQSFLR  
 VFKGIDSIVDGEVMKFVNMSMIKNNFNFRHVKSVHHILQCCNVYWQAPCAVFLNLYY  
 KSLWIIQDICLPYCMIEYQDNPA MGILPSEWLKMHFQTLWTNFKAACLDRGVLTGCE  
 LKIVHRDMFCDFDFTDAGSNGLMAPFKMQVRIARAMMVVPKSIKIKNRIIFSNTAGSE  
 AVQSGFVKPTGTRDITYVAGPYMKFLNSLHRLFPDTKTAALYLWHKISQTNKTPVLK  
 DVPDDELAELVSYVKTNSLA FEETNVLDVVPDSLMSYARIKLN GAILRACGQIQFYAT  
 TLHCLTPVLQTI DAEEYPHVLGSAAIATPVAYLAEIRGRTALT VQTARQPVAATGRL  
 RPVITVPMVNVKTYGVNGNNNVFHCNGLGYFAGRGVDRNLWPESSPFKKTGVSAMLRK  
 RHVMMTPIIDRLIKRAAGQTI STFEAESVKRSVQALLEDKDNPNLLKS VILELIRHLG  
 KGCQDLSSSEDVQYYLGDYCMILTDEVLF TLDNIAQSGVPWTIEDAGALIEDRQDADDLQ  
 FVDSDDIATASCQPPEEQLP TPSAGALLAGKKRKINALLSDDL"

SEQ ID NO 10 CDS 9468..11528  
 /note="transport protein; ORF 7; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 7"  
 SEQ ID NO 11 /translation="MARELAALYAQLSALAVDLSLVIFADPRSIDGARILKTKTQIEN  
 LNRDLLPLLREQNSVETSSLSLEVEHLAKNIEDKLGELERSLRQRYSSREHFETLHLR  
 PECHYHSTVTFQFYGGGLIDVNMCLINDVELLCKRLGVSFYCIGANEALSGLNRLVLT  
 LSTLRGISPIPHPDLYVTSVPCVQCLREIELVPNQGSLLAVLADRHCDHLCKKVRÆ  
 PIHGLFETELSQLGLKVTKRSDATQHGVRSADQLRESSLA AIQDHNIFKRV SASIME  
 LSNLIYWNAGQTGLQTGTENEC SQMARLLTHEADMHEHRLITPKLSATHFYDCFRPD  
 PIESLFCGGLFNSIDDTINALSRDCSVTFFQQANYTNVMRKQNELFTRLNSILRQGS  
 GSQKPATPSEPRTTTVAATAASDV I KDAQYRKEQYMKKVARDGFKKLT ECLQTQSAVL  
 ANALCMRVWGGVAYGEASELVNHFLRRRFVALPWEARCRSDQILFENSKYIKNSLYS  
 QRLSREHVEIITLQFYGLITGPLTRQSDLFPGPANVALAQCFEAAGMLPHHKMLVSEM  
 IWPQIQPKDWIDQTFNRFYQLPEGDLNAVQKSAWCFIRELVLSVALYNRTWEKTLRIF  
 SLAREKLSISNLDVKGLTSGLYLTYEQDAPLV LISQNTGWIFKDLYALLYHHLQLSDG  
 HDDN"

SEQ ID NO 12 CDS 11515..14004  
 /note="glycoprotein B; ORF 8; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 8"  
 SEQ ID NO 13 /translation="MMITNRTRLLRAWV VIIAIGTAVGENVTTPKGATTTAKPTPGP  
 STPTPPENPPRAEAFKFRVCSASATGELFRFNLEKTCPGTEDKTHQEGILMVFKKNIV

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PHIFKVRYYRKVATSVTVYRGWTETAVTGKQEVIRPVPQYEINHMDTTYQCFSSMRVN  
 VNGIVNTYTDRTFTNQTVFLQPVGLTDNIQRYFSQPVLTYTPGWFPGIYRVRTTVNC  
 EIVDMIARSAEPYSYFVTALGDTVEVSPFCHNDSTCSVAEKTENGLGARVLNTYTI  
 VATRQPTTETRVFADSGEYTVSWKAEDPKSAVCALTLWKTFFPRAIQTTHEASYHFVAN  
 DVTATFTSPLSQVTNFTGTYPCLNDVIQKTLNATIKKLSDTHTATNGSEQYYETEGGLF  
 LLWQPLTPLSLADEMRELNGTTPAPPTTTSTANRVRRSVGTNEQATDDLAAPQLQFAY  
 DKLRASINKVLEELSRAWCREQVRDITYMWYELSKINPTSVMTAIYGRPVSASFVGD  
 SVTDCVAVDQASVSIHKSRLTSTPGICYSRPPVTFRFLNSTTLFKGQLGPRNEIILTD  
 NQVEACKETCEHYFIASNVYTYKYDYVFKKINTSEISTLGTFIALNLSFIENIDFRV  
 IELYSRAEKKLSGSVFDIETMFREYNYTQRLAGLREDLNDTIDLRDLRLDLSEIV  
 ADLGDVGRVTNVVASSVITLFGSIVSGFINFIKSPFGGMLMILVIVAVVLIVFALNRR  
 TNAIAQAPIRMIYDPIDKMQPSGGKVDQEQIKNILAGMHQLQQEERRRLDEQQRSAPS  
 LFRRASDGLKRRFRGYKPLENEEAQEYEMSK"

SEQ ID NO 14 CDS 14122..17166  
 /note="DNA polymerase; ORF 9; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 9"  
 SEQ ID NO 15 /translation="MDFFNPYLGPGRPRPHSHRGTDAPAPAGAGAVQPPPDVCRLIPA  
 CLRTPGAGGMIPVTIPFPPTYFENGARGDVLLANERSMWTARDRKPVPADPDQDSITF  
 HAYDVVETTYAADRCAEVPSRFQTDIIPSGTVLKLGRTEGTSVCNVFRQOVYFYA  
 KVPAGINVTHILQQALKNTAGRAACGFSTRRVNKRILKTYDVAEHPVTEITLSSGSML  
 STLSDRLVACGCEVFESNVDAVRRFVLDHGFSTFGWYSCARATPRLAARDARTALEFD  
 CSWEDLSVQADRSDWPPYRIVAFDIECTGEAGFPCATRDGDAVIQISCVFYTTREGAP  
 NPPNILFSVGTCDPIPDTDVLEFPSEYDMLVSFFAMIRDFEVDFTGYNISNFDLPYL  
 ITRASQVYNLRLNEYTKIKTGSIFEVHEPRGGGGGFMRSVSKIKIAGIVPIDMYQVCR  
 EKLSDYKLDTVARQCLGGKKEDVSYKDIPPLFRSGPGGRKAVGSYCVMDSVLVM  
 LKMFMIHVEISEIAKLAKIQARRVLTGQQLRVFSCLEAAARENFILPVPTPEGQGG  
 YQGATVINPIPGFYDEPVLVDFASLYPSIIQAHNLCYSTMIHGRDLHLHPNLTPDDY  
 ETFVLSGGPVHFVKKHKRESLLGRLLTVWLEKRRAIRRTLAAACDDPSLKTILDKQQLA  
 IKVTCNAVYGTGVASGLLPCINIAETVTLRGRMTLEMSKSYVEALTTEDLRTRLGRE  
 VTARHGARFRVYGDTSFLFIACDGYSAEAVSAFCDDLAARITADLFPPPIKLEAEKT  
 FKCLLLLTKKRYIGVLLNDKMMKGVDLIRKTACKFVQERCRAILDVLHDPEVKAAA  
 RLLCKRPPHAVYEEGLPAGFIKIVEVLNASYLDLRNSVVPFIEQLTFSTELSRPVC  
 DYKTTNLPPLAVYQKLASRCEELPQVHDRIPIYVFDAPGSLKSDLAEHDPYVRQHQIPVAV  
 DLYFDKLVHGAANILQCLFGNNADTTVAILYNFLNVPYKLF"

SEQ ID NO 16 CDS 17261..18511  
 /note="unknown; ORF 10; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 10"  
 SEQ ID NO 17 /translation="MLVNELSVVLGDWEVTFHGRFSFVNLTRLQTFKGHGGYARVRL  
 PFSLDQLLHQHFAFGLVTRLKELPPFSDCVALIAPLDSGGDADAARVAPGFVLDSSRP  
 LTVWVNASGRHTIRFCLLFLKPIDLERAVTYVFGENGGARSEGTPKPTCATESLPGGP  
 LRVSGEASQTSPhSFVAYFPTANSVACLSSLRLQVRPFSDDAHRDARISPKYVTF  
 SN SGGNVCKASVHTLSPSRCKTAQMEIIYAPGDPNAEIVLGQSGPVLPHTTGGRV  
 LGVYA DAEKTIQPGSSAEVRVQLIFQOGAAARGDLAFLVTGVAPEPLFVVTPALL  
 LSGCTTHL RLFNPNGTPTTIKRDTLVAAAAPCPVVRSSADDAPRDLVASPD  
 TGALSINAFTIPVG FPGVVSACHVSLRDNGVHERMNH"

SEQ ID NO 18 CDS 18520..19749  
 /note="unknown; ORF 11; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 11"  
 SEQ ID NO 19 /translation="MGTPVRFFRGEWQTSSLVDNGTPRYSSLVWAATIH  
 DGYLTLVNR SELCVTERSPLCPACPSIGRLVGKRFPGFAFASATLGDGRGTRTVFYA  
 FGHDRDNPDI PAVVERADRELVLRVHAPQTTTRVSRYGLKVFAIVTVVRPPGVFL  
 HFPPQDRVPIALTD ACSQEGSRLTSEEPWIKIQGFPVLSDETAHPFLLTQKTKP  
 FTERKFCRLIMDNDQRSA VNTVYLGKQHVVRVTVPETIVTDGPVTATLSLTGNAP  
 IAFRHNPFYFELPWSSTTAI FTPVVYVGLTVCIPPNCSEKVFVRYGNTYVSAFNR  
 KLTAIISNHAHNGGFRIQDCEWPPN"

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REIEILVTNVSQAPVYISTGTQLGQAI FVFAPRFGGPAKLRQLLGHRSRALPGGVT  
VDSQKLCRFETMYLFST"

SEQ ID NO 20 CDS complement (19921..20544)  
/note="R2; similar to IL-6"

SEQ ID NO 21 /translation="MFPVWFVLFYLS CWAASPTLAPPPTAAGINVLPQWAGNRASLDR  
TRGRLSEVGLNIQRWFVYLCHHSTLCRVREYPRIMSFVHFPI LMSNVECQRREFRGAE  
CMNAMVRGLRAYESYLTRLRMLLDDAPGDADAAAIGSAVTVVL SALSLEELP VNNK  
IGGAESNEKTVRALGGQSPRDVVL SAFRILEYLQMFLRDGRRAIAMM"

SEQ ID NO 22 CDS complement (20777..21778)  
/note="thymidylate synthase; ORF 70; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 70"

SEQ ID NO 23 /translation="MIVLVHLGICYVKKIIPVCVAGIAAARLRVFS APEGAAAVRCAC  
RGDHGELQYLAHLDLIIKHGVQREDRTGVGTRSVFGLQARYNLRDEFPLLTTRKRVFWR  
GVVEELLWFIRGSTDSTELSRRGVKIWDAGSRAFLAAQGFGRREGDLGPVYGFQWR  
HFGAEYRGADANYEGQGV DQLRYVVDLINRRPHDRRIVMCAWNPADLARMALPPCHVL  
CQFYVARGELSCQLYQRSADMGLGV PFNIASYALLTYLIAHVTGLTPGDFVHTLGDAH  
VYNNHVDPLLLQLRRTPRPFPR LKILRKVARLEDFTRADLSLEGYDPHPHIEMEMAV"

SEQ ID NO 24 CDS complement (22245..22592)  
/note="R3; similar to Kaposi's sarcoma-associated  
herpesvirus K4 viral MIP"

SEQ ID NO 25 /translation="MRGLFVCVFFAVFACVVDYAFPMGSM SGPAPELCCLGYVTHLPP  
PGLVVSYSHTSSQCSVDAVILNTRRGKLCANPGDDAVK KLLQAVDKRPKKGRTRRS  
LIDDSEEGLGSGI"

SEQ ID NO 26 CDS 26846..27409  
/note="Bcl-2 homolog; ORF 16; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 16"

SEQ ID NO 27 /translation="MAAVQGPPPPPEEENENSLPVDVYAIEGIFLYCGLGQAEY LHHP  
VFSPIKEFISAF LKDSARLYERLLRHTDYRSLRGLNAIGQGM LQINTDGRHNWGRALA  
VLGLGAYVVDKVKDDERLLTFAIAVL PVIYAYEALESQWFRSHGEWEGLRNYCERILRH  
RRNARRHMCYGVAAAGLLALVALFAIRR"

SEQ ID NO 28 CDS complement (27515..29125)  
/note="capsid protein; ORF 17; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 17"

SEQ ID NO 29 /translation="MTPVYVGGYVDVVS LPKIEKELYLEPSIVATLLPYTDPLPINIE  
HVPEAHVGHTIGLFQVTHGIFCLGKLTSHD FLALASRLAGDSRAAQIQLNPMRPDPLL  
EMLHTWLPELSLSSLHPEELQDPNHPPAFQHVS L CALGRRRGSIAVYGPDP T WVVS KF  
DSL TREEAGKITVNC L D L C E R Q V T P P E F A A P L E T L M A K A I D A G F I R D R T D L L K T D K G V  
ARVARSTYLKASQFP CAQHCGNRDTRTMSALPEDNITIPKSTFLTMVQSSLDNMRNQG  
HRTYVSAPPSMPATAAYPSWIPPELTVPSYAPPVAPFPFQSAFAPQPSPYAATYYS  
PTYGYAPAPSRHQKRKRDVELSDEPVFPGE E VGIHKDVMALSKNILD I QADLRDLKRA  
ASQTSGAQDADQRPQPPPVQFSWPQTYASAPYLAYQPQWYSGTDTHLHAPQPYQSAQG  
IQQTQPPPPQASHHAGLATQPATPAPAAQESVMSNAIP SASAPRAGACPLDPECGQ  
SARAPVEASAQPAPVSQIQKMFCEELLK"

SEQ ID NO 30 CDS 28998..29897  
/note="unknown; ORF 18; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 18"

SEQ ID NO 31 /translation="MFIGRGSVYGSRVATIEGSKYSSFSIFGR LTTSTYPPTYTGVM L  
GRCLREP KEMSAGLRGLMWRVIRCENLNTFLPGELRFLHLVLC E M Y N G L N V Y L L K E A  
IANTGTRDDIVLGRKVPVEFWKIVYDGLKEMGVSDATLLSETKRGALWLYFN GRPCLL  
KGLGDYVFCQLGLSHSVRVPENLTDGNYLYNLG SVIPCRLLVALSYCLAFWGHADHE  
PWVRLFAGKIFILYLIISGHIMPRKSILEQVGTSGYGGFVEAVCRDVRVAVHGIPAWDF  
ASAAPALTSQQT D Y L F A F N N S V V"

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SEQ ID NO 32 CDS complement(29905..31548)  
 /note="tegument protein; ORF 19; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 19"  
 SEQ ID NO 33 /translation="MRTSEKCCMRYPKPARQITATFWAPHPNNVLFHKKPSLIEERR  
 NAFVMRNQQLALRVHTLRKNLLRLELDNVLQTHQRETEVVMRDLETIQNMVGDLSRPG  
 RETANAQTSLNPPQPKIAPQTHGDAFVVTIAPGDPGFTVNQDLRELLPSLYMNQNWQL  
 PQYGPWYSSLTDNAMQRRVFPRLDLRGTTNFQNSTSLKLMSAVISTAASITQDFYADVR  
 NVSDTQAALCLLNGYYCHRTGTPLPPTRNGLWDNLGTLATLVSHLKQNTKGLGFET  
 YSNPRORASLAPLNKETKYNAFFTNHVIYATLAQSGLLPKSKNPGTGQPPGPDVYI  
 LATTLFSEDVPPFQAYQWNLRLAGLSALGCLVLVYVLELAQITPRSPHRRNLASLLG  
 GRFSKVEDPSGSKQYLKKGQLFDLTENYISPILSRAPDAPTSFLFPGAYLAALAEAKA  
 ISHLKHTRPFVNLTGSRFNEIFDILNQKLTFRDAGSLIQAQTSRLTAEEGLAILSH  
 PSPPGLAHEIMKSQFGVYDDYDRVYFLVLGYLPVATSVV"  
 SEQ ID NO 34 CDS complement(31043..32095)  
 /note="unknown; ORF 20; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 20"  
 SEQ ID NO 35 /translation="MAFANQCKHVATLEALPASRKRAGTRAHLAVYRRLIKHRSLLDI  
 LKFLSIRPTLRATKNVKFRIFFEVSLGRRRIADCVLTNSEHQKTCYVIELKTCLSAAV  
 FPGNAIKISQRWQGLHQLTDSVAYIGRAAPRGHENWSVRPWWLLFKNQKTLKTIHTESS  
 AFPPTFINTTSAALNGFFSQWEDAHVRKMLYEIPKTSAAANYRNLFGPPSKQRSVYSQ  
 TISDRRKKKRVCDAKSTAGAKGSHAAKPPAPARTRQRAANAPTGNRSGHARPRNNSKH  
 GRGSAVPGQGNRQCPNITKPATQNRPADTWRRVRCHNSPRRPGIHGKPGSPSGAPAKP  
 VHEPKPMAATIRAVVQ"  
 SEQ ID NO 36 CDS 32094..33767  
 /note="thymidine kinase; ORF 21; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 21"  
 SEQ ID NO 37 /translation="MAEGGSGFGDELVRQMRDRKPRWDESSDDTDDVDTESTDLEYDD  
 VFPVVDTHGLMSPGSQNYDVPTSPSGTPWELLHPDALYAHPRCPPKRAVVPGGGARPK  
 VSAFSARLQYVGRQSFQDRETRQLTGAQFSSESEHEYAEIPERTTTTPVESGDKRNF  
 SGRGGAISGPSSTKPSHGAGLTRKTKTSLSVSLKNLLRIKDDDAKVDVPRPVTVPVHL  
 MQPHPMTEYRNAFLIYLEGVMGVGKTTLLNSMTGMVPOENVLSCEPMKFWTCVYSNC  
 LKEQRSIVKQGTGHLITSARVYACQSKFALPFRATAAGIGRNLQPWLVGNGSTKPN  
 WIVFDRHLLSATVVFPLVHVKNRLTPDHLFQILSLFSAHDGDVVLTLTNSSEAHRR  
 IQSRGRKEEKGITQNYLRQVAVAYHAFVCTWVMMQYLTPEQMVLQCVQTVSIEDICNM  
 NSRLTHRFLTTLKLHEQSMIPMAEMLVSVKEHVTLMVCLGLFKELRKLQILIVDAG  
 EHLDDACGLWGNIIYGQVMSNEAIKPRAVNWPALESYIQTTLTKLEGNGAY"  
 SEQ ID NO 38 CDS 33754..35868  
 /note="glycoprotein H; ORF 22; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 22"  
 SEQ ID NO 39 /translation="MARISFIFFTIIRCSVTDKYVYDEKSNVELEFNGTIYQINWRN  
 VSKELTSIVMEDAWYDSLLLEPLSVTLEKRKSLLRSSIVNVHNNDYTFCKSSSDHVIN  
 LTVDFNYSSLPGFTGNFNVMTHALTQGVLLTKRELFTNSTNIMDLFYAEKINAEMFKI  
 TFDYSNVIISGIITENWILVSVTNSSVKSNNMQCVALLFGVPSTFPALKGYVSYRDLV  
 VKNSNYALGVIAPKSYNTLDLAFLPKNFTFMFVSVIDSPLNAIDYLGKLLAEAKGA  
 CQNPSNENDILSFFFEVTAVNFLFIKNLQKQQLVNVGCVVRHVAALSLMHLRLCYP  
 TFKLYELNLETLSHIAESQVFNLPANSMLSLSVNDQEVVFSMFKIVYNTPKVGGKILN  
 EIVYITNYMYTKYSENYQLTNTFRNVNMNMYEVLTTIKLVNTDSSVFYPYILFTSMCN  
 NVEISYMINQIAKPDITIFRVFSPCFLSLRFDLDENKLRSAPQTSKRTGSELAQGA  
 SGFWRLHAFHATRINEFSVINCTRLAWQVTALMPLTNITYVISSVRPDHARVYEV  
 EVFLNSAMFVSAVYPNCSHFTPPGTALHIPILYNFSAPRIGPLCDSIVLSYDENQGL  
 QTM MYVS NPTVQANLFS PYS PF DNDNFH IYHLWLMNNGTVVEIRGLYRRHALSAIAL  
 VF AFIGTMSALYFLFKLFSILA"



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SEQ ID NO 40 CDS complement (35865..37073)  
 /note="unknown; ORF 23; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 23"  
 SEQ ID NO 41 /translation="MIKISDLKARLVGGAVQLSNGEYVCHVVYSSALAAMVGLPGPAV  
 PLPLLFKKFSTIYSNMMPLYAPKRPELSMLRIMVSPHPYALNSCLCVGTDEGERGVSL  
 FRDPVIRSSDFEDTPITVNSKLVIASNSLFLHCRPFSPATVKTPPVTLTNNKQITIN  
 ELANTTQEYDPNAPPTLCSALPPDNKKLRSILKQPPATSESNVQSDCLLADIFFAMGS  
 RQPQIGESPITAFNTVTIMQRANNSIMFLPNLKLKPIQHLFLKHVLLQRLGLENILFH  
 FKMLYANTCKAAGPYQREYFESMLSRVKQRLEDMVFCLSIESHDFQKDFRVLSRAPQ  
 RLLTATDKYFLMFPQNRRLAIQVGAEVIESICDGTPLSEVLANLSPRVTIQKETGNN  
 LLKFYALLTV"

SEQ ID NO 42 CDS complement (37123..39321)  
 /note="unknown; ORF 24; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 24"  
 SEQ ID NO 43 /translation="MLLQGPVLLPACPATVAANAPSPANSDFKTQLAIFCCLATNNEI  
 LENVSLEALDRAMQTETTFYACRALRRLLVLGEGLYPFIHRQGGIVGKTGNEYAGPGLI  
 IDDAIGCTFSHIETHFLPTVFTYELSDTVLVQSDERILRSLYCSPLMVCVNYQSMF  
 RILCRYLQIWEFEECFAAFTRTLPEHLIGTCYQNYFKLLEPFKILTARCPPPCAKLH  
 LNYLKFNILGFTSDWISHPELHRVQTVIIHNIESNPVLLKNLSKQNKQDIKVASELI  
 IDYQNIQVNSLDVNLQVKINKKDPGKKPYKVVVVTPKSTYYLTFPEVPPIFRVAMCMS  
 VAEHVCHSCDRLYPNTTEFLGPGETPRVLEAMFSRIQYAPKDRDYNFIFNADQNPDRYE  
 QARHDHQTEPLPDMFDPVKHMSLHNFKISVFNTNMVINTKITCWSLAGTFESIIDI  
 LTNNFVMKKFSVKEPSFTVSFVSDNLCNGAAINVNISGDMHLHFMFAMGNLRCFLPVK  
 HIFPVSIAWNSTLDLHLENQYIVRRGRDVFWTNTNPSVSSKDGCVNSWFKAATA  
 TISKIYGRPLLKKLSDELNPILSVPYARIDQVKNTIFTTLETRNKAQIQTLHKRFIEC  
 LVECCSFLRLDLGALNRAARLGTDFDSKRIISHTKSKHECAILGYKKCNLIPKIYVRS  
 KKIRLDELGRNANFMSFIATTGHAFSNLKPQVIRHTIRRLGLHWRHKAKI"

SEQ ID NO 44 CDS 39323..43459  
 /note="major capsid protein; ORF 25; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 25"  
 SEQ ID NO 45 /translation="MEAALEVRPPPYMATEANLLRQMKESAASGLFKSFQLLLLGKDAR  
 EGGVQFEGLLGVYTNVIOFVKFLETSLAVACVNTEFKDLKRMTDGKIQFKVSVPTIAY  
 GDGRRPTKQKQYIIMKACNKHHIGAEIELSTDDIELLFIDRETPLDYTEYAGAVKITIT  
 ASLQFGVDALERGLVDTVLNVKLRSAPPMFILKTLSDPVYTERGLKKAVKSDMVSMFK  
 SYLMDNSFFLDKSDIAVKGKQYVLSVLSMDMVGAVCHETVFKGTNTYLSASGEPIAGVM  
 ETTENVMRKLLNMLGQVDGGMSPASYANYVVRGENLVTAVTYGRVMRTFDQFMKRI  
 DRPNAQPSVDDDRDAVADGQDSLAKTPIAAAVIQIGDKLVALESQORMYNETQFPFPL  
 NRRMHYTYFFPIGLHMPRPQYSTSATIKGVEHPAEQSVETWIVNKNVLLSFNYQNAL  
 KSICHPRMHNPMPCGQALGQAFDPGHVHRYGQRSEHPNMNLYGLVYNYQGNVAH  
 VPDVALKATMTTDELLHPTSHETLRLEVHPMFDFVHQPGAQAAAYRATHRTMVGNI  
 QPLAPNEFQNSRGLQFDRAAAVAHVLDQSTMEIIQDTAFDTSYPLLVCYVIECLIHGQE  
 DKFLINSPLIALTIETYWNNAGKLAFINSFPMLRFICVHLGNGSISKDVYAHYRKVFG  
 ELVVLLQQALSKEIAGHEVVGRRPASELINCLQDPNLLPPFAYNDVFTNLLRQSSRHPMV  
 LIGDEGYETENDRDTYINVRGKMEDLVGDMVNIYETRNNADHDGRHVLVDVGPFNENEQ  
 HMAVLEKLFYVVLPACTNGHVCGMGVDFDNVALALTYNGPVFADVVPDDEILDHLE  
 NGTLREMLEASDIHPTVDMIRTLCTSF LTCPFVTQASRVVVTQORDPAQLLTTHDDGRYV  
 SQTVLVNGFAAFIAIDRSRDVAETMFYPVPFTKLYSDPLVAATLHPLVANYVTRLPAQ  
 RVPVAFNVPPALMAEYEEWHKSPMLAYANTCPMTPTSLSTLASHMHMKLSAPGFICHAK  
 HKIHPGFAMTAVRTDEVLAENLLFSARASTSMFLGQPSVMRREVRAVTFEVNHELA  
 SLDMALGYSSTITPAHVAAITSDMGVHCQDMFLMFPGDSYQDRTLNDLVYKQKAGCQRF  
 GPGQIREPVAYVAGVPHSDNIPGLSHGQLATCEIVLTPVTADVITYFQTPNSPRGRAS  
 CVISCDAYNNESAERLLFDHSIPDSAYEYRTTVNPWASQQGSLGDVLYNSTSRQVAVP  
 GMYSPCRQFFHKDAILRNRLGLNTLVTEYAARLTGTPATSATDLQYVVVNGTDVFLEQ  
 PCQFLQEAFTPTLAASHRSLLEDEYMSNKLTHAPVHMGHYMIEEVAPMKRLLIKGNKVAY

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SEQ ID NO 46 CDS 43491..44408  
/note="capsid protein; ORF 26; similar to Kaposi's sarcoma-associated herpesvirus ORF 26"

SEQ ID NO 47 /translation="MALDKSIVVSVTSRLFADEIANLQSKIGCILPLRDAHRLQNIQA  
LGLGNLCSRDSAVDFIQAYHYLDKCTLAIVLEEVGPNSLRLTRIDPMDNYQIKNAYQPA  
FHWDNYSSELVVIPPVFGKRDATVSLESNGFDVVFPAVVPEPLAQTVLQKLLLYNIYYR  
VAETTPTDVNLAEVTLYTTNITYMGRNYALDVPVGSSSAMRMLDDLSIYLCVLSALI  
PRGCVRLTSLVRHNKHELVEIFEGVVPPEVQALDLNNVSVADDITRMGALITYLRSL  
SSIFNLGRRFHVYAFSSDTNTASCWCAYN"

SEQ ID NO 48 CDS 44433..45242  
/note="unknown; ORF 27; similar to Kaposi's sarcoma-associated herpesvirus ORF 27"

SEQ ID NO 49 /translation="MSIPKIMTVSRDNEGTVCEVAVDNGRHRAMIYYPKTTNLANERA  
DVVKEAFDTETPVDIVKQIVNEGLAISKKNCVRLALYLYFYLYQYVCFALLLTWQLNPY  
MDPPGLVFAVNPMPGPKHVTKLPHPAIVAVGCGADAICKNCSVPDIKTELGMVYHNGSS  
DSGQRAHYGLALLKAAWLVMGNVCPEPVVRQGAALLGPWNRTEWSDFKSAMAATTFCC  
SRGVLWSPiHEKNLCRPTWNDVINTSVFTNESLCPNIPVVPESVIVLNGDA"

SEQ ID NO 50 CDS 45408..45683  
/note="unknown; ORF 28; similar to Kaposi's sarcoma-associated herpesvirus ORF 28"

SEQ ID NO 51 /translation="MTAHTNGVLTGTFSTSQPESVQVSPFYRVITKPPVMGLFFCVA  
MCVIALVWYVMRRVCCCKGRVVADSCRDPQPAYEMLNVRLRPHGTNP"

SEQ ID NO 52 CDS complement(45733..46779)  
/note="unknown; ORF 29b; similar to Kaposi's sarcoma-associated herpesvirus ORF 29b"

SEQ ID NO 53 /translation="MLQKDAKLIFISSNSSDKSTSFLNLKDAHEKMLNVVNYVCPD  
HKDDFNLQDTPVACPCYRLHIPAYITIDETVRSTTNLFLEGAFSTELMGDAATSAQSM  
HKIVSDSSLSQLDLCRVKSTSQDIQAMKPCPLHVIYIDPAYTNNTDASGTGIGAVIAVN  
HKVIKICILLGVEHFFLRDLTGTAAYQIASCAAALIRAIIVTLHPQITHVNVAVEGNSSQ  
DAGVAIATVLNEICSVPLSFLHHVDKNTLIRSPIYMLGPEKAKAFESFIYALNSGTFS  
ASQTVVSHTIKLSFDPVAYLIDQIKAIRCIPLKGGHTYCAKQKTMSSDDVLVAAMAH  
YMATNDKFVFKSLE"

SEQ ID NO 54 CDS 46905..47135  
/note="unknown; ORF 30; similar to Kaposi's sarcoma-associated herpesvirus ORF 30"

SEQ ID NO 55 /translation="MENDTPKDKISEADFQCCQAFFHRPIRDLISSGADALNHFSLSLSE  
SDGHKLERIVLLDLVGTECLSYTTIAAKNVK"

SEQ ID NO 56 CDS 47093..47746  
/note="unknown; ORF 31; similar to Kaposi's sarcoma-associated herpesvirus ORF 31"

SEQ ID NO 57 /translation="MSLLYHDRCKEQMTRVNSPICRFHNVSNLQCLDCKRYHVCDG  
GRNCVIVYTRENLVCDLTGNCVLDNVQDVCSYGPPERVPDAFIDPLVSHGTRECLKS  
DILRYFETVGKSEAYSTVVKNQQLNGIIGRLIDATFNECLPVMSDGEGGRDLAASIY  
IHIIISIIYSTKTVYDNLFLKCTRNNKYDHIVKTIRAQWMMRVSTGDPSPRSATGCFT"

SEQ ID NO 58 CDS 47683..49077  
/note="unknown; ORF 32; similar to Kaposi's sarcoma-associated herpesvirus ORF 32"

SEQ ID NO 59 /translation="MDAHLNRRSVAGQCDGLFHVILPRGFILANNITCGERQRFPAH  
TWFAASGRSTKTLVWGRVFQNTDPGRGDGPGSPWSGLAISLPLFTTNGKFHPFDVVI

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LKADTPDSGSSWTVKFLYMSLIAAYRNAMRGLKDKVSQCTDAAVDGEVHPLTVLKEAL  
 VSPDTATRPVSACNPLQMLTGLLQSRVRDDYVTHHRALERPGNVRGQVIAPTRTEMPN  
 GSPSRVRLGFRPPKQANYPKTWAQARHVFSRRAYYVCVYDNEELDTKWQRQDPRPLPL  
 DWSDPVAYLLEGDLFLGAKQNAFVDSIEKTCRCQNYTIKQFFPVLINRDNETVDLIKE  
 HFIEACFVIRNQVSERSAWVKAALFRNDSNTYWKDVLGLWEHGPHKLGTAIKLPTSEP  
 CNADVNWSWLLCDEDITRSISGQSTVCLVVSPTLTAWLVLPGGFVIKGRYDLSSSEDLN  
 FVASRYGHPASSHS"

SEQ ID NO 60	CDS	49049..50059
		/note="unknown; ORF 33; similar to Kaposi's sarcoma-associated herpesvirus ORF 33"
SEQ ID NO 61		/translation="MATQRHILKSFLNKECIWLRHPGTSFVRVYTATTAHSAVFDP PVTSENAmslNfLNMIVIMKPKFEGPCVTVYMNKDILDFCATESVAIRDVPGRADLC LIRFGTlSNAPRSVPIPGPLNPHPRETVPGlTKQEIYTSQTVPRGQIPDAIKGKEFH QINPFLWFDGGAFWQLFLSVDFMLLCPALDTPVSLARIVGLLTQCDKSTCKICTGAHV HVNPyRGYTPPDSQGTSPSCPLISCGARRAADVLVTGHVNLGLLFDPKASPKVTKL RLKRNP RPVP IEDAMSGVTAEGTEVQPTSLPWALIRLPDLASRVMLYGCQNLKSI CLR SY"
SEQ ID NO 62	CDS	complement(49977..50960)
		/note="unknown; ORF 29a; similar to Kaposi's sarcoma-associated herpesvirus ORF 29a"
SEQ ID NO 63		/translation="MLLTSYRERLQNNLRVVTDGGCENWFRQPPVVISGNDKTERMAH PCLGVIHAVNAYSSVLDDYLQTYRRVQEPMPAPTLGKPRISSHATLPRLTEELTNYLK QTCCRVQMANAKDQYMEYQSAQRTHEAFLECPVYAE LRQFLANLSSFLNGSYVPGVCC LEFFQQQLIMHTFYFIASIKAPEKTHQLFATFKQHFGLFETDDVLQTFKQKASVFVI PRRHGKTWIVVAIISVLLSSVENVHVGYVAHQKHVANAVFSEVIATLSRWFPKLNLI KKENG TIVYASPGRRPSSLMCATCFNKNVSR CFLSSGSRIASRDWLN PAGE"
SEQ ID NO 64	CDS	50959..51942
		/note="unknown; ORF 34; similar to Kaposi's sarcoma-associated herpesvirus ORF 34"
SEQ ID NO 65		/translation="MFPSSFLNNGHPETERRFVKGVLALDLCNDTPGQFKLVETPLN SFLLVSNVLPESRPVRDCPQEGFD FEHIHL PKLTRMQRVLG RYCDHVNNDDTCVNVK ASSNSQGALFYLPYGQDEWNWALT LRKDKLVKMAVEGLSNPTTWKGLEPVDPLPLIW LLFYGSR SFCREPECLYERNFGMKGPILLPPHMYAPQKQVMTFVHHVIKYVKFLYVNA GGGLETEPSPPFASRLRAAIARLGDVEADDA YLSAKCMLCHLYKQNDTISI HETHVG GVIALGGDGARYITSSVRAQRCTSRGDFVLIPLYNIEGLVSMIREHGLGSS"
SEQ ID NO 66	CDS	51923..52372
		/note="unknown; ORF 35; similar to Kaposi's sarcoma-associated herpesvirus ORF 35"
SEQ ID NO 67		/translation="MASAAKKMLIKSELESEINKKLSISVDFRFGADSAVFNAQYKG TRESLRSYNSLKKKDDLATVVGTLTSLREKQSELGLLKGFNRKKIEF DAVADAVRD LKDELYGELEILGTLDNESVPVEEESP KDDIIRWKLERLPRVCPKSP"
SEQ ID NO 68	CDS	52278..53585
		/note="kinase; ORF 36; similar to Kaposi's sarcoma-associated herpesvirus ORF 36"
SEQ ID NO 69		/translation="MNLFPWKKSPPRTTLLGGNWSVCPECAPKALDPIPKVQTDVDR T ASSHITVIKTRKTIAQLKIPNNWGC SHQATDWTAVLGRGSYGVRSM SLGRVCVKHFG SRREFFYECIFNDIVRACREKHPLNRGGDRILCFLEPCVPCRALIFPQLTGNLLNADL KHVNPERLAVEFSELREGVSFLNNICGIVHCDISPENILIKGELTTAYGR LMIGDLGS ASLHTGTPWTGVMVTSKLG FVQHTYHFKAPARFICKHIYRPSCLLYRCLLSCAGGPQA HMLNQPFQITPQLGLTIDISSLGYSLLACLEKYLQPADPFPQOGALADASSES AHPLF YLR CMVPRVIAE IFSVAWDVPLDLGIDSSGHAPAIPLREAYRRFFANQCSLYRAQYK EDALENASSRLCNSK LKLV LQKLLVRDYFSHCNCGDHGFFLR"

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SEQ ID NO 70 CDS 53566..55008  
 /note="alkaline exonuclease; ORF 37; similar to Kaposi's sarcoma-associated herpesvirus ORF 37"

SEQ ID NO 71 /translation="MDFFSDEPMVQEMALLDIDEQORLLSKMSLANFLKHERVRAFFS  
 DNKKEISMPAIRFVYNFYLFKAVGDFIGNTDVYDFYVTCVFRGRLTRLSEVYDACLN  
 MHPHDRHHVICALIEQVTRGQNINPLWDALRDGISSSKFHWAIKQONSSKKIFNPWPI  
 VNNHFVAGPLAFGLRCEEVVKILATLLHPGEAHCENYGFQMSPNLGVFGVSLDFGIN  
 VRSDPKDGLFHPDCKIYEIKCRFKYTFSKMECDPIYAAAYAKLYQKPSMOTLKGFlys  
 ISKPAIEFVGEDRLPSESDYLVAYDKWEVCPRKRRRLTAVHHLVKKCMIHNSTAPSD  
 VYILSDPQETGGQINIKAHLSANLFINVRHPYQQVLLQSLVQVEYISLSKGTKNLGT  
 QKNFIATGFFRKRQFQDPSCCTIGEFAPLDPHVEIPTLLIVTPVYFSPVAKHQLVKQA  
 TEFWAASAREAFPELPWDLSSLCANAPPTP"

SEQ ID NO 72 CDS 54963..55172  
 /note="unknown; ORF 38; similar to Kaposi's sarcoma-associated herpesvirus ORF 38"

SEQ ID NO 73 /translation="MGFILSVCKRPTNTVDVKGEPIDVSKFDPPIIGESIVLLTADG  
 TAPAALYKPKTKPSKHKNKLSDFV"

SEQ ID NO 74 CDS complement(55255..56391)  
 /note="glycoprotein M; ORF 39; similar to Kaposi's sarcoma-associated herpesvirus ORF 39"

SEQ ID NO 75 /translation="MKISRSDSFILSSWVKLLVILGLMFIMSAVVPLTATFPGLGFP  
 YFNTLVNYSALNLTVRSSAKHLTPTLFLFLEAPEMFVYISWAFVLDGYLLCYAWAILAI  
 FKAKRVHATTMTSLQTWIVLIGSHSVVFMSSILRLWTIQLFIHVLVSYKHILLASFYCI  
 HFCLSFTHVQAMISCNSATWSLRVLEQQIPENSLDLTLRYGKPIGANLYLSLIAMEM  
 LVFSLGTMMAGNSFYMLVSDIVFGSINLFFVLTIAWYINTEFLVKYLKHQIGFYVG  
 VFVSYLILLPLPVVRYDKVFISASLHKVIAVNISMIPITCILAILRIIRNDWKWCAKS  
 PEYAPLPQGPKEKTTKVKYSPELNALYETEEDVSDYEDAYPKYI"

SEQ ID NO 76 CDS 56526..57932  
 /note="helicase/primase; ORF 40; similar to Kaposi's sarcoma-associated herpesvirus ORF 40"

SEQ ID NO 77 /translation="MNAREVALTGHVLHISLHSTHEREKLIWQVHLLVCQQCGIQGD  
 AAYLFVTETLSNTDWNIPAINRHAPSINEHGRNYMQWELRTRLRNPIIQLLSRQPGA  
 VNVRVSEPNMVIIVGCERALDHSCSVRVGTGAYLHCDTTMDFSLDSVVSPTREFWFSEMF  
 SHCLVSNIEVYLKTTGGLYRASSATQCRKRAKDGA LGILDIFNCESREIQVAGQKYT  
 LSIATATFHVLWVDEACMWNGALAEFFRALHNKLFGRGVAPTLTIVCPGATPEGTP  
 FPPYFSAPPHLPLVFGRRRLDVTAVQELPKAQIAVHWPPFKDSILGDQLLIPGISPK  
 KPGTVPVRWPLWVEDVNLSCETTESVARIVDPHSIVIIFSSLLCQHLKCHRAVFN  
 ELEYIATICSSDLRLFIQEEYNRLLATIFTWAAASGYTAAIDKTTVFIIKAPQLSAAV  
 SGFCPSLNSCRRKQCYEG"

SEQ ID NO 78 CDS 57917..58528  
 /note="helicase/primase; ORF 41; similar to Kaposi's sarcoma-associated herpesvirus ORF 41"

SEQ ID NO 79 /translation="MLRRLKITVHFLSQEQQKVTRLEAHLGLPVQETSHPPDWLKCE  
 VCSASVFLKIPAGVLYAGLARDPTREAKRDSWLDCLVEGATLLLNNSVLPICALAGIL  
 PTLFANRRVCVNFLLPRAWVKSAPICPLPIDCVTPPQFVVTKRGPICWYKEWPLPVD  
 VDFMYYLQEALCVFSVVSNGEGTESHADNIRQLEKFEKVLCLF"

SEQ ID NO 80 CDS complement(58525..59343)  
 /note="unknown; ORF 42; similar to Kaposi's sarcoma-associated herpesvirus ORF 42"

SEQ ID NO 81 /translation="MDQILKRLMGEGHRSEAVMPETECSSRGYPYNYVFPRLMLEVHK  
 KNSICMASNTPKLCVRGRLNVPDLGVHVRTRLQSATFTGFVFACVVEHEDMIDALDIY

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PHVFSDRVQLFKPASASVTELCCILSMLENYDKPPLSFILSALDRARYLHERYTCNDS  
AFVLYGIEVIASSTLAAYHELNPPQGILRVPLVRFKLHKLLDENADDMKGLLKPIYLE  
SFRLTENVGEEEGHAETFNIFYCGTIFTRHLHNASVLKYFQITSLHSIPRQTLF"

SEQ ID NO 82 CDS complement (59297..61027)  
/note="capsid protein; ORF 43; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 43"  
SEQ ID NO 83 /translation="MFKMNPFGSGTCLVHPTELSISLFEILQKAYVRGQTLHSSLR  
NPGIFGRQLFIHLYKTALGSCYDNLKDWTNFETTLKTRWRGVEHLTPFEKRSTFES  
WARTVRLTVDQLLLNTINQVLHTRTVLSYERYVDWVVALGLVPIVRRTPDGDTIARIQ  
AHCQOMRKTYASGDVTISRIVDKLAQEITSIMTDVTSIYIPDYAEVSVEFNGDKAAYL  
GTYRQKDITVEVVSRIIYNGRVAFDSPLYRLFTAIMTCHRATAEHAKLCQLLNTAPLK  
ALVGSTCNDMYKDILARLEQSSQKTDPKRELLNLLIKLAENKTVSGITDVEDFVTDV  
SQNIVDKNKLFGTGTESTTQGLRKQVSNTVFKCLTNQINEQFDTISNLEKERDDYVKK  
IQCIETQLLOSLPEGGRPRHDINILTQNTLQALSGLRDPNTINLSECHIPKGSSVNSF  
FSQYVPPFMEMLKELTSLWEGEMFQTYNLTPVVDNQGQRTSIAYSQDQTVSILLGPFTY  
IIAKLTHMDLINHSLISLSLHDIADQLYVDSRSLSVYINDIGHKYCEQISQPGTDGPNT  
EASNGGAAP I"

SEQ ID NO 84 CDS 60966..63338  
/note="helicase/primase; ORF 44; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 44"  
SEQ ID NO 85 /translation="MESSVGWTKHVEPNPGFILNMTSDAKVRGVVDHVSRLSNITTSP  
PEMGWYDLAFDPAEDSGPFLPFTVYLITGTAGAGKSTSISALYQNLNCLITGATTIAA  
QNLRRRLKTFCTIFSAFGFKSRHINIAVRKAHQTAGVSIIEQIQQELSKYWPVIVDI  
MKEVMAKKPNGMYGTISNANFETLSRMTGSPCLWTSNIIVIDEAGTLSSYILTTVVFFY  
WFLNSWLNTPLYRQGAIVCIVGSPQTQNAFQSTYNHGTQKTEISSCENILTFMIGK  
KVVSEYVHLERNWALFINNKRCTDLQFGHLLKILEYNLPIPEVMSYVDRFVVPKSKI  
MDPLEYIGWTRLFLSHSEVKAYLTNLHTCLTLGGDTRDTKLFTCPVVCVVFVKPFEEY  
KRAVNLTNLTVEWVTKNLFKLSNYSQFVDQDMSIVATESTERSTQVTFITKFVKNSH  
VSLNGKTKKICGFGQTYFEFKRILDSELFVETHSQDRPEYVYGFLNTLLYNAMYSFH  
AYGVTRSHEKYLQDLKFAPLPAALATGRVDLQTVREELNLEDDIFYHVCSPPPPAGIT  
SLQVLVDITYCALKDVFASRIKVACRWFGGEFEKETFSFTVNMVVRDGVDFVSPSERL  
NGLLAFASSTVESYKIKGYTFLPVAFGRQCGLPLSDDLKMKPSLVVQDSSGFIACLEN  
NITKLTETMEDGSVFQVCCAGDYGVSSNLAMTIVKAQGMSLERVAVVFGSHKNVQTS  
VYVAISRANVSNYLVMDSNPLKTLREPVNDNTSAKHIVRALHNPNTTLY"

SEQ ID NO 86 CDS complement (63379..64437)  
/note="unknown; ORF 45; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 45"  
SEQ ID NO 87 /translation="MAMFLSDPSRTPPATPRMLPIPGAPRKKRTRRFLFAGSRTGLPV  
PPGYGGPPVIDMTAPEDVFDQDSPPTTPKTPDETDSHSENSDYSDVDEEDEPPVSSPP  
RIDPHARDGESFNQSGRLPTVITSTGATTTPPSAPAPLTAFGGPRPVAVVTGQHRAPQS  
SESDSEDDFFIDYEDTDES GGEADGFS PRASPAWSGDISRSPAEGGWSSDEEPPVVA  
GSTAGQETIIISDDDEVDDRGSVETWDES DADEGTGATDVIDLCSSSDSDDDADHVT  
GGVRAACKRRASRRDCNGDDVDVIYVGTQGPKRMTSTTGGGATSNPEGPGVSGRQTM  
AATPPVCGNDNYPWPWLD"

SEQ ID NO 88 CDS complement (64479..65246)  
/note="uracil DNA glucosidase; ORF 46; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 46"  
SEQ ID NO 89 /translation="MEGWLKTIWWSKMSPEVLEEPSTQTLTLLSDSWLEFLNLSPLFKQ  
KLAALLKRVMDMSNVTVIYPPIDRIMWWSYCCPEPEDIKVILQDPYHRGQATGLAFS  
VAPDYSIPPSLKNIFKEIANTVPGFTAPSHGCLDCWAKRGVLLNLTILTVERGKAGSH  
ANLGWDWFTSYIIISLSAKLQRCVFMWGRKAIDKATLINGQRHLVLKARHPSPLATA  
HAATGSPWPQFLGCNHFKLANDYLVQNRRAVDWNIN"

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SEQ ID NO 90 CDS complement (65222..65731)  
 /note="glycoprotein L; ORF 47; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 47"  
 SEQ ID NO 91 /translation="MRSMYTSLSLFITCGFFLITCCTGLVVPCCCKIIPLSDFIFPEPF  
 EIASFHLTNLALCPGLCTATLRYKADRSTTEICVNGFHLRAFFIRILYKLNYSVPREE  
 LQLLNYMQYSLDEFLAEFEDFHINGSSEGTAYTRPPLDSDRSTKVSRIKRVITRRG  
 DLWRVGLKQ"

SEQ ID NO 92 CDS complement (65999..67168)  
 /note="unknown; ORF 48; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 48"  
 SEQ ID NO 93 /translation="MAVSIPVKGVNRETESNWSIVTTFERHGNADRAIRSLRFFFKG  
 VDHPGFLASLVILKDVITIDSEKTIERTDLIPLLQGVRFVTOQIYMHLKDHASESPMAE  
 IWRDCKERFCLALELACGCQRCASAAARQLRACQACRPPKLNPHKQOCVAARLLTAVY  
 NQMVLRTVSVSEFCLNALMCVPREFGFVSGDVRVETSRVASCLNLSWLYLILDSYVR  
 TDLTNLEMAMSRACRIHGLSAGDPFYSALVWLKNSYACDTNTFFFTVNSTSVTTPILM  
 DICASLTGPVPDVIKINMLPLVNDQMHPSVCVERANFTGSCPKVSPTHLDGLKLETT  
 SLTLAADSLDDILOALELICGDDEGILDSYISDINTETEVDESSIEEEIVFEELS"

SEQ ID NO 94 CDS complement (67398..68303)  
 /note="unknown; ORF 49; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 49"  
 SEQ ID NO 95 /translation="MSRHYGKDHLLNHMYKFHYPPGLMIVGEMNTLTVNARNPLYQAA  
 TLRVERALYLSKILQVLMQHRQGERFIVPQCRSNMVYCLKELHKITNDRIRGLINSVL  
 PLVDAGCVGFDEELVRVLPEILKLEYPHAHELLPPHDPTSPLSWCLSHMVGVTKTFKG  
 EVKEMIDTFHDLSPVPSFYQLASLVKKFFLVEEVIYEDYQDTQFNVLNLCFFWTTVIK  
 MYQSCIFKDKLLDTIKACIELLKGEARQFFGWYDLNTPNLGSSALVKYTEHLIRALSV  
 DSSAIPIGEICSHLHHCKHALLNLE"

SEQ ID NO 96 CDS 68494..70038  
 /note="transactivator; ORF 50; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 50"  
 SEQ ID NO 97 /translation="MECASLGPISGLIADLNLNLFCLYGRSRVKTRGAATCNVPCAE  
 CAQGIVRILTERALCCTEKMFIASACSGVVIPPOLARVLHDVYAEMKAKCLGAWRRLI  
 CCRPIMAIADSVLVTYNTLDAEGKLELRLKALCKLVFQPIFLQRIAPMQLLANGKM  
 VPDNYFTITGTAEKRRPVVTGSTSGMTCPGSSLPVDSLILPVCEPGLLPAPLVDSLNV  
 LENPEIILSAPPLSQFVITNTHPSLPQSUSIITPTQGVVPGQCFMDTWKAVSQSIHQ  
 AQTPIILAAALTGSTSAAPGPHIACSPVAGTSRQVEGSAGVDGCKPACVPQPALPPNP  
 AKRMETVAQLGNAPVKNVHIGGRVYAPLVNIPIDLTSPSGSGQSPADIANTPESRMA  
 AGSPPPFAETAATVPAKRKQPREVDADKRLKGDVRGAATVNHPFPGPSGMRVREQGLFD  
 LIESSTDVTANASGPKNDDMLAAILQDLYGLQSPPAIDSPSSNSDNEEIFPEVSPPS  
 SGHGSP"

SEQ ID NO 98 CDS 70355..70888  
 /note="R4"  
 SEQ ID NO 99 /translation="MPRVKTQPKRPQVLEFMPLDLHGHTHEMDSQNLCPDGDLLGS  
 YIYTENNGPFSQIMHNGQSNLTGTGESFGSYAAGDGLGGSVSGMYGNNTGEGACSKRP  
 SACRKRSAALIHAASEASVAEQGTSQGAHAVSDRIGRDGGADNRLKVSARLSDKTKS  
 ALRSHPCRLRCYSLMFNT"

SEQ ID NO 100 CDS 71468..72160  
 /note="R5"  
 SEQ ID NO 101 /translation="MGFGNIRLGRWLCFMVWVAXIARGRSVCPTWHLTDGKYEAVYRH  
 YLEECRKHEGSGSLDGSQTKGSGTKATTEANISIRPNVVTSGQNKEPPGTAPRAESS  
 HDLPRIKQVNALRLSTPELAQPLPVVKSTPRESQSGGTPWNARPHAFIMHTNDMLNPS



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VVLSFRAIRARSTRDTEQSVDRNTVTTTSYRTPGRPSLFQARPSSHGARLPPSPRTMA  
RYAESRTICDQN"

SEQ ID NO 102 CDS complement(72401..72820)  
/note="unknown; ORF 52; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 52"

SEQ ID NO 103 /translation="MSSTRPKTRAPKKELTMEELAAQVQKLSVENKQLKKLINSGDPT  
RSGSDPVISNTEKEAKIAAAVSALCNVATRKIEAKVRAATAKAVTRGQMEDALAGISI  
RVDVSMDETTTRGGIAASADGALRRRRAQSRTRNNAD"

SEQ ID NO 104 CDS complement(72884..73198)  
/note="unknown; ORF 53; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 53"

SEQ ID NO 105 /translation="MTGSIVLALALLACLYLCLPVCATVTTSSSTTGTGTPPVTTNPSA  
APSVTPSFYDYDCSADTYQPVLSFSSIWAVINSVLVAVATFLYLTYMCFKFVETVA  
HE"

SEQ ID NO 106 CDS 73274..74146  
/note="dUTPase; ORF 54; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 54"

SEQ ID NO 107 /translation="MAEVTAAHTVPYAFDSCKFEEIPKNNSSRIALRNKFPVVVKPGEP  
LVVPLGLKIIRAPQCAFFLSGAPTDEVYHTGLIDQGYRGEIKLIVLNKTKQVVTLYR  
GEVNVSLIAFMYASPGPLKCPILNLPHYSLDAGFDVTS PHAMTIPPTDRTPFTLSLYY  
KSPQLSTPHVPLIVGRSGLATKGLTVDATKWTQSLVHLRFYNFTKEPIDIPANSRICQ  
VVFIEDHVP SGWNILRSRVQLGSTLQISWAKIRFTDVATLPKTHPLNSRHTQSQTEP  
ETARGAKGLGSSGL"

□

SEQ ID NO 108 CDS complement(74207..74839)  
/note="unknown; ORF 55; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 55"

SEQ ID NO 109 /translation="MAAPGSFWTCCGFSPFGRVGCQYRPLPDPLNECPHWRTEIAMG  
LPPGVDMGDVQKQAE MCTAALRQTYLLAVQSNKITEYLRRFDAARVPAGCQETVRIQIS  
KLKSIQNV IWNAMLSLAIGDITVDES AFHALLNKRADET VSLLEMEKLATTIASDDSV  
TWAAEINNVLVDTEASSNP SHPVIRQPTPQLAVADNIVPDKIIQDAQADG"

SEQ ID NO 110 CDS 74851..77337  
/note="DNA replication protein; ORF 56; similar to  
Kaposi's sarcoma-associated herpesvirus ORF 56"

SEQ ID NO 111 /translation="MVDEIRAIFSTSGDMAEVIDILTETQATASFFCVLHDRGDAPI  
NTPHAVIKLCLPAKRPGGPRCLPLMVNLPAWQVNLFLTGDAPLTS DNIKDRIDLAQ  
TEEILEPILSVLACKRSAQQT KHD SFKSKVAFRAKFVSALRKVYKMTSPYWMITLL  
GSFEASFVL AGTFYFFQSHICTAETLVHLTRLFSSSQGQSLVTNTYDELGRVFGSD  
FLGIVPNFWAYLKYKMQDDVESRAIDQTINSIRGGLMLS PQDLVHFIYLSFYECMNA  
QTFLSYSRTTSSLPTPATVNPQLCRRLEADFKHEVMAYYNKASYLSTYITILTVPAP  
LPDGYENFQELACQYWCQSRDVAEIMTRINDQYPQLNLTKDLSGLLDLAALDQYSGG  
PKENLFTVASRIPTYRCEFLNKQYFVLMHADCIDAYWKQNIIVPEDAQLOGLTDQDLT  
SRIFYCDLGLSLPTFKQQILVSRHEYFNRLPVYRWVLD FDLKVTEGRRTLNDIYNIC  
VTLRQVILETLQLIGPLKPNHPVYFFKSACPAVTWPDDISDTAFCHCDAKIGMRIVTP  
FPSGYCLVGSAPLVSLTDILNRVVKLDTRLASEYPGILEDKGPFDSGIYAKGRCVRVP  
HCYKVGPGGELSRLKIIICHPEESDKSAYLKNFAKVS NLLHAPGDSVTKNHGLVYA  
ITDENEGFLESKTKNNLPKTTITDLAEKIERTEKPLIDWAATAVWPKLHDTIQRFFPD  
DRIGQFASVSFMHSGDNIIQVKPQKGNFFCINH KHRNHTQTVRVFLTLHSTKESEVT  
VTFMSQCFAAKCNHNSPTAHFSFMVPITGT"

SEQ ID NO 112 CDS 77578..78906  
/note="immediate-early gene product; ORF 57; similar to

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Kaposi's sarcoma-associated herpesvirus ORF 57"  
 SEQ ID NO 113 /translation="MRYVFHALICFIGGSSSDFDDSSSDEMDDLSTPEPEPSTTPN  
 SFPEGPKSQVVALPKIRKRSRSETPVKIEHRSPLNRSRSRSTRSGSGQRSNQSGRYV  
 KRFKPTVDAPRHREPWHRGKGKAPFIRRDAMAGRGRRTYGHYRGKAALTRSIKESI  
 KKMHL PSTML SRAHDKKVFEGLLPRHLGQCFQVCLPAPPPLQPEVFTDRQLTAIVKSG  
 GRRDALVAKKVSLAKLTSLYKPLLTFTVTRNNQAHWLATRKNLTASAGLEALAAFIIE  
 GLAWAQVCVSNRSLNDSNLDIILDSSQSVCWTFISKIRHLHIQCFLNQGEVSLVKQ  
 LTYLVCINNRLAEANLAGEVKLNFKLGMLIGFALTLPALLAEHLKSGESLYLFRSFL  
 EKYRPGDVMGLLNSIVVEHYTKCRSAECVITTHAMVSGGENNKGLEFFFPV"  
 SEQ ID NO 114 CDS complement(79266..80513)  
 /note="R6; similar to Kaposi's sarcoma-associated  
 herpesvirus vIRF K9"  
 SEQ ID NO 115 /translation="MATWRPPQSGGPSAMGLREWIVTHANLATYSGLFWADDEKTRVV  
 LATTPWSVGFYLDYLRDGMKMYEDYCNQRNIPPLPSGRSRLGQAKARLLGAIRKSAYFIEE  
 KDVLRRSFSFANVVFRLRSDEEMLCRLCPRASGVAAELRGLRFRMFKRKGADEAGRVS  
 EYTVKQLLGLLRTRPAGTFTMTAPATEASATATASGEDGRQDNSQGGAVALPGEHALP  
 LSASSGLSACLAPSVDDPWGFMHIQVYYYGVQLAQTFTHSGMGVRLSTRPTDKNEHHV  
 CMAPGPLQLWLPAPYMDDDFMLSRLVNALHALEDGIVLCSCQYGIMMNGYGFNLWF  
 RGNTSNTSEPRRVPSPGVGHRVFDTDYMLKLAQSPRPSDPGPPDPFAQIWSAWSLYE  
 EEDQSQAPICIVVHQREIYRHF"  
 SEQ ID NO 116 CDS complement(80686..81933)  
 /note="R7; similar to Kaposi's sarcoma-associated  
 herpesvirus vIRF K9"  
 SEQ ID NO 117 /translation="MAGRGVDIRAWLVAAVESGEYRGLVWENEDKTVVVPWNKVTAD  
 RSVWNSEKFFDDYCNMRGICQGEKPSHYGRFRKMRFLYDMRHHKSIRELKFINKAYGR  
 PGARYRLFRLLPEPVVSCAMCNLMSSSTETQCLGLISEFQYDQGGSGRERRRVFTATV  
 LARSMDKNKRVREHRLPGAQLTFLYFGETVGLERVHAGIRVCSRPCPVLAGHACCF  
 QDERTLFLPSPGVVDCQFAREDLRVMHKKCEKGVLTITLTDTGICVKNLENREMKVLTN  
 NEEYKDLPSRQPVQVDFMDVYLRALARSPNPGDEPPRDYALCLSVQSPNPVDAP  
 IAMRLRYVCETSSVCGTEGCFYPGTTVTSEGRDSCSFQMEDPGEGETSQSHDPAVELGD  
 SGPDSMDDPDAGTSGEDDGVACS"  
 SEQ ID NO 118 CDS complement(82262..83317)  
 /note="R8; similar to Kaposi's sarcoma-associated  
 herpesvirus vIRF K9"  
 SEQ ID NO 119 /translation="MERPVVRTKPSLRGWLVECCETGRHPGMRWVDEERTLIRIPWN  
 HDRGSRGVEEAENKIFIDYCRSGILHAAGRELTAKECKNWLSAIRHSQTVSDVSTK  
 DNLSTPYPDRCRIIRLLPITVRSCARCDQASGTTAMLRGLREEAVNKFGPVGAGVRYT  
 GAVGAGGEQCWMLRIMFYGGDRVGEVVTESSNGIRVLPLSERRPQGHICAAPIAEQ  
 LVPEIPGHLAEFQAEALRFLDKDLLRGLAFWADPSGIYIRWLGHSLAFVQGNVESTGA  
 VAVLSCANACRAFNLVDYMTAMARTSPDGAALPQACVYLYFGGVPTPEGGVQSTVPLI  
 IQLWHECLWRALSAANV"  
 SEQ ID NO 120 CDS complement(83491..84252)  
 /note="R9; vIRF"  
 SEQ ID NO 121 /translation="MDSGCIYACILDENSEGIINYLEQVCGIGLEPGMPLPAPLPTLV  
 PTRSAYARAHRLGVPEAPLPHQIVPFWRLRIQVYFGVLALDHTSQDRRGVRLHPRPV  
 PHPGHLCFYGTGFTVWFPSPDREKLTAEQITQIKTMLVAYNEGIYVHGNETGVYVDNR  
 NRETLAAGNDCNGDIQREVMFLSKQKIFNFMGMFRKLARSPPGESHAPCNGATLYL  
 SQQPGAQESPVQVPIVSVVCQDELVQGHMNP SRWCS"  
 SEQ ID NO 122 CDS complement(85052..86209)  
 /note="R10; similar to Kaposi's sarcoma-associated  
 herpesvirus vIRF K9"  
 SEQ ID NO 123 /translation="MAAGESRRGPSRYGMALKEWLTFKADSGLYPGLFWADEQKTRLV

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LAATPPSPFNNDYQRDGHQHYDAYCELRHIPLPSGRSRLCQARGRLLAGVRKSKYFEED  
 KEFPTDQFPFTALVFRLRSSEEMSCPVCPRVCALRLELRNMRFAMLRGMLHALSGPS  
 VSDQERRYREGHQDGHDAQDDDAAYSSGLLRARLMACAAPSAGDPWGHMHKIYYGQ  
 LQAEELLTATGQIRLSSKPTNKAGHHVCVLDGPLQAWFPPIQTTESVVRLEDAK  
 WLVDGIIFCSTSRGIMFTITGGPNVWFQGNTEPYSLPHRAYTGMHVWAFDTRDYLDD  
 MARSPSPRDTGPPAAFVKLWVSGCSLGEERNSSRAPLSIIVYQTEIYRHF"

SEQ ID NO 124<sup>b</sup> CDS complement (86355..87527)  
 /note="R11; similar to Kaposi's sarcoma-associated  
 herpesvirus vIRF K9"  
 SEQ ID NO 125 /translation="MAERDMDLKAWFIEAVESKRYPGVEWDDDKTIIRVPWNRCTDS  
 RVDEDYNKIFDDFCARGVCQTGSHAQFKKIRMLYAVRSHRYLRELTPPSKAGGVSG  
 ERYRLFQLLPEVTVGCDLCNLIATTSLSHSCSMGSCVREDVFERTRRPRAKAPLRVSVY  
 KRKSKRLQDSSAQPVLGAVEVSFFYFGENVGVOILRAGSGVRICGLPDPKRPGLHCCA  
 DNPLTCFLPSSQLIPCEFARADLQALQKTCERGLICVMTESGICVKNLEERNMTALTN  
 YSENYIELRPSQPLQAFDLLHYLRELARSPTPGDVPPRDCAWIFMCPSTQSENTWDAP  
 IALKLRYVCNDDVSDDVSNGAAGDDSGDEGPSGAGVGASGTTGSTSSTLAPYGRK"

SEQ ID NO 126 CDS complement (87894..88961)  
 /note="R12; vIRF"  
 SEQ ID NO 127 /translation="MAEGRAGSIRVNRPSGLRAWLLDCCDNDKHPGMHWLDEEKTIVR  
 LPWNHLKGAGGVSDDERNMYLDYCQFKGIRQTGNRRLSVRECKNWLASAIRHSQTVED  
 VSTEENLSAPAPNRCRVIRLLPIFVRSCLCNEADATGGMLLDVRNEVTARFRYLGA  
 MEYEGAVGGDGEQCWMLRLVYYYYGRLVGNMEVGS PNGVRLLPAPKRPLOGHVCAGIR  
 PEQALLPHTPQDMFPHQTSMLKWLKKEIIRGLMIYADGSGIYIRYMGHVPAFLLGNGG  
 SLEPVDIINNARVLRVFSLAQYLSAVSATPPHGTFRFPAAYASLHLGGVPTPEGEPCPT  
 IPLSIQIWHECLWRACGDAQ"

SEQ ID NO 128 CDS complement (89122..90216)  
 /note="R13; vIRF"  
 SEQ ID NO 129 /translation="MTEIETHNHLRRWIIISNLEANTFPEHLWCDEEKRSFRISWHR  
 GMSGMQPVVAYCLDRDLECGRQHNVSCEKRLRLRVLENAGFEQDDARATTTFRGGER  
 FFYLRPAVDPLCYACILDSSHSETVLNYLEAACVHGLEPGTLPPLPAPAEADGAARSVY  
 ARAARLATVAPPHPDQITPFWRRLRIRVFYFGSLVAEHTSQDRRGVRLHKRQDPKPGHE  
 CFYGTAYKMWLPKPQLDGPLTPEQRETVCIEIINGCEGVFLHGNELGMYVDNRTRHTV  
 RCAGNDAEGNHAQRAVRSSVKSQIFYVMGELLRLRLARSPVPGDTVPSNAVTLYLGGRRPG  
 SSKRPQVPVTLVICQDELTHGDIRAARWIL"

SEQ ID NO 130 CDS complement (90462..91544)  
 /note="unknown; ORF 58; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 58"  
 SEQ ID NO 131 /translation="MGTYTSEASLAWLSFMSGTVSASPFIICFIYHSLYFVEPLISVE  
 NIIFSWGAVGLHGLLLFCIFGLPAWLSRQVDVPCTISAFILITAGSMASLTGLVDLPWV  
 HVSIFVGSCLCLLL CVVAANDVVYLCPTIAHRYVELGFLAAFSVYYFLVLKNLFLAPV  
 FLLPLVAFIVGGVCSLRALRSHPLYEAGLQRRHAIFSLTSRRYITYSIKQALEVCGWD  
 FYLVTVLIGGAAAGTLSVGLTTPLLGLVHYFFVFHVGLFCCLGLVFRSNVLALVYVL  
 AAVLLTLTHVLGPGTHNLFTRVCFVTVFLLTMFGAIGCELQIIRKKLQRAANSRIV  
 LGVCACGNLLMAVVFFSLNKVELGAL"

SEQ ID NO 132 CDS complement (91555..92739)  
 /note="DNA replication protein; ORF 59; similar to  
 Kaposi's sarcoma-associated herpesvirus ORF 59"  
 SEQ ID NO 133 /translation="MPVSFHYGARVDVDALGSISRVDHIKGIKKGVIIQISGQGRAP  
 VLSVLSSVG DAGVLGLRLKNALAPLMVYSDMTDEVSFSTRNTSLGNTFTHTREMFGVN  
 IAEMNVAFYHHGDES DAEGKPQFVRTTIAYGDNHTSTVHKSVVDEPNLPSFHDRLEQA  
 GTGNRLFLT VKLTLLKWL RQQKTRAKQVTVSLSETLAVATFTVDGVSKI IDFKPD  
 TPDAKWTCARGRKL DVGVSDDLTHVSLESLSVAALNACKIPGFFLPGRWHANEILE

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VEGLPLTDSLADVRLGVMLLKVDPTDRNNAVPGNLSEGADPEGVPELPSPPRTPDLDL  
KEQCVPIAEDGAEPDGGAKSLRTSGSRPEKKHGKRKHSSSPSRGKGKTKTPRATFNP  
LF"

SEQ ID NO 134 CDS complement (92868..93812)  
/note="small ribonucleotide reductase; ORF 60; similar to  
Kaposi's sarcoma-associated herpesvirus ORF 60"  
SEQ ID NO 135 /translation="MFGLSIVTAAMESPDRFLYASDHPGFLALTQETWQNRWFPSQIS  
LHEDSDEVRLLSPTDREFYQFLFTFLGMAESLVNFIENIEDLVKEFSNHDVTHYYAEQVA  
MENIHGKVYANILNLF FGGNRGDLMIYAKKIVEDATLAKKIDWLHSRVRKATTRAEKV  
LLFLVIEGIYFISSFYSIGLFRLRGIMRGVCLANDYISRDELLHTRAASLLYNTMISR  
DESPSVAYIHGLFREAVEIETLFIIRSKSRDVTMVNVGDIEQFLQATADRILKSINIPP  
LFGARPPNACPLSYTSAKSVNFFERDNSEYVTSVHNDL"

SEQ ID NO 136 CDS complement (93794..96160)  
/note="large ribonucleotide reductase; ORF 61; similar to  
Kaposi's sarcoma-associated herpesvirus ORF 61"  
SEQ ID NO 137 /translation="MNTETSFSAAKSAKPLTLVTDAGTGGCSSSLDPERCAESLVNSL  
KATLGWDIEANSLTGLLWHRIMEDRCLVTVRDYLA VFGERLSDEVRAFMSKHEAALDG  
LLQDFKQSKAYTNFVNCGYLSAVRFYDITYVLRTQGSSPIFESVAQMFMRAVAVFVACQC  
IKFPCLRETLRHLVESETELD EMYLVGYAFHYISSQIVCCATPVLRSAGLRGGQLSSC  
FILKPSMATEDKTLKALHEEMSPLLASKSGVGIDVSSFAEHKNITSCLKLINAHVGYP  
NDNNIRPVGASAYMELWHHQICDFLNAKMPENQERCHNLFQGVCPVPELFFRLYETNPD  
GQWHLFAPEVAPNLLKLYGAEFEIEYNRLVAAGKHSSSLPLKSMYALINTVIKTGSP  
YVLLKEALNKHHCETQGSAINCSNLCAEIVQQPEGQASVCNLANISLPKCLRPHRGE  
SGVEPGKGDVTFGFELDDAVEAAVIVNACILGGTAPTESVRRGQEERSMGIGVQGL  
ADVFAELGFGYLD AESA KL DVEI FOAMYFTAVQTSHEIVLLGEGT PFRGWERSRLAQQ  
VFHWQTWDGVKPSHPPLERWEQLGRSIAQH GIFNSQFLALMPTAGTSQLTGYTEAFYP  
FFANIAASKVTSKEEILKPNVTFFKRVKPGDLRTVRRYGGDVASFPEPLKDRYKIFLTA  
FDYCPKQLERAGARAPFVDQSQSLNFFLKEEQATRASYIRDLLLTGYRLGLKTMLYY  
CRIQKQTKLNALQCLDQVVGDNISSEGAESNCVQKADGERTKVCLACQ"

SEQ ID NO 138 CDS complement (96163..97158)  
/note="assembly/DNA maturation protein; ORF 62; similar to  
Kaposi's sarcoma-associated herpesvirus ORF 62"  
SEQ ID NO 139 /translation="MKTRDANVNKLNDSLMRLLPPPPHRVSLSRGRDFSKGVRDLLSK  
YVVSTTTGVEAIKDGFLSVSPTCQTYGDFLIYSQTMSSQEPRTYLF SFKQTD TGSSI  
DMLFTPTSLARLSRMDADSAPQTNRIACVWYGHEGLLDAIPNFEELETGSLHQFLA  
PVGPLVQTVHSTFVTKVTSALKGNVVAREPVVTHIGLTLPSDMFVDLDDSCPSSLRDE  
PLPAHSSIIYVCLTYIRVNNRPALGLGFFKSGKGYCEIAAQLRDFYSGVIRTKYIQLQN  
DLYINRLAFGVVCR LGSVPSGLQPSFQSLHFKGAALPVLKFTEFVS NPGSWKLFL"

SEQ ID NO 140 CDS 97157..99976  
/note="tegument protein; ORF 63; similar to Kaposi's  
sarcoma-associated herpesvirus ORF 63"  
SEQ ID NO 141 /translation="MASSIPAARADNGDENTGGLYKLTDNLLTCTGSLQQLKLLMEFQ  
LKPLPTAHL L SMPTVTRFLNTAFKIDNPLVSFIQKHPVFFLMRVARLPEPVITDHQSA  
ETSTGILSEVVNLNTAIRKPHESPA AKNDYLDNRILAMITEYIHHVTSRTPSGIP  
PTPPMGISHLPCVEQILHETHRQYWNLTLPESLFDIGEVASPLQTWLILSYCKKLQL  
APPPPLFPVDELARRLV TGHHELFVPLSTSLETYITMPVSKQRAFEIYSVFAKSKNIV  
DGTPILAFTDTELTFTPELLFLYDFVIESLCKNQAYGCSRNAIEHF IKKGIDFMAEL  
GAFIEKTCGYRSTVSLSNVRAVKARLASGLSKEACEDFRAMILMTPHETTPKWENFT  
DFLEMVNQLTLYGYFYFECLNQYSPTSISLAKIQNINLRVDAEQSDRALWRTPLIGSF  
PPPWKLNNVLAFKFPSTPVATLQKIYKAIPSYLMRSLFEIAANKSWGNIALAESAPLT  
DIQTAEPDQGPVSAQVIAKYCSRLQISATDYDAAIVSSPGFAAEFIKTKLYPILSEVL  
RNTSKKNRSLFQIRWLIVFAAEDARDLAPIRRSLALAYFOIMDILEEKHSPEFYNLL  
DYLQETFR CIRQVIEPATCPQEFLQYLFTFQNIPIAASFIQTSMTFVDDLKNGIPGIL

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DLVSLGAAFYNMKLLYDSTLDTVEIPTTEEGQPIVVSFVFKSTIRVLEKLLQEAVIAL  
 TQTSEPMYAAHIRLMQHLTYMQKIAGHEIMTTQLPSVFHEIHEGYLQCFKRFKRLMLH  
 VTGSCCYSLTRYFGFLYQPLIPDTIVQKILNFNDKTDTTDDILKSLSQPVRQGPLSA  
 ENESSRSLSKNNVELLQKLYDDFRTASTNNNPTSILKEYSGNYNETQVSVDWSTYNLV  
 TYTAPDDTLKFTPVNTEALDRMFAE"

SEQ ID NO 142

CDS

99980..107626

/note="tegument protein; ORF 64; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 64"

SEQ ID NO 143

/translation="MELPPIFSKFIEGVATTHQADCRFGQYAGSQCLSNCVIYLAQS  
 YFNRESPVTDNDLDDVLRQGATLDFILRRSGTLGYNQYLAQLHHIPSFIKTNEWTAAI  
 FQSQEYFGLIGLDAAIREPFIESLSILTRNYAGTVQYFLFCGDKAGAVIIKNKTFY  
 LFDPHCVPHVNPSPAHVISSSDPTAILEYVSPDREYTGSLYIMPSEYVNPHEYITN  
 HYRTITFAKVHGPIDISTGIEPCTIEDIPSPRSPDVTSKSSNLARVPRTTDTSSA  
 KPPPATLSGLRGAEPPTSYDPATNDADTKLLTPAPAQTAVDHPEFQTPGATLLLSE  
 LSASRGRKRKLSLQRYSDSDEASSDDEGAPRRRVHDDAISAEVIWMDDDISPLYSPS  
 ATPSFDDVFDSPPMSPPEFTYEDATEDTDGAFLEQIARDAETPFSAFDDDLITDHDFFSL  
 DKKIEQLIKYEAPSOHLNPNISDKQNGRAVREAAALQAMDKIMINIILEHGLITDAQAR  
 GPSACKNVLQFFILWGEKLNIPISDAKQVLELDLQLIPLHTAISEGKFKQGAFKKHLT  
 TKINRCLASMRATHADAQKKLASAFNVEGSQISSSEAKISVRALKEQIANHLSPGFLA  
 VYSADEVKHLRDKIQDLKTGIEQRNKEIQQEELFFDAMLTALDTFQPPPKTAFPMEIF  
 PHRKTEVMLDHLASITTRLTDATEALNNYLETPPDQGTHITNIPNFSSIVANIISTL  
 KILTYAENDMQLNVTPMATYRRQLLYLGGELATIFNLEWPYETVPPVQELPLVARAKA  
 KMESVTKMEKNQQAALDQILGDAETLLDTITATSGDENPVRAMSIPILETYITNAGALI  
 GSSRNQRFEKLKAAIHDLASSESFIIMLLNNTRLDNISDNLAKIDGILTNNTRFLSNA  
 TVSKTLQTLGGSLIRECVEALNKRSPSSLNNARLLAVQTLGHASVPDHEHTLTRIVSG  
 VASAQKESAGDDPDRWTRVTGHLNELKLVTTSQSRVDKATRRKLLMIITRDLKEAEVSQ  
 ETVLETRWQENVLKFQPSTSKIEIDFLQSAPSAKARKFAEKHLRTLITQFNHERPPS  
 EATAVPMDYTPPTIPTPQAVSTATAEKGKAAWNKIQQAFQDFNFHLIDASDWQEMASE  
 YSRHGSSLPGTVGPKLVRFMESISNTLDDILTQKLASLLPNGPAFRPPAFDWIAPYQT  
 RVNAFLKTIGLPMVRNLADKIHHCQCTVSHAVQSADLQOATVGTSLERPAAEYCRILS  
 DMQVAFNDHGI AVRSEAAAYTDAINSPANVVT PPKPNLEAPKKLITATDALTVEDFPD  
 FLKTSILQQEQRILALQRAEFQOLEASISAAERLRQSTRDEIAGKMATAITQLLPRAP  
 VAISSRPLNLSKPIDFLSSTVYDKILDKEPYETAIAGFAWLEIATKSMVYSQQNETQ  
 QLNVLLESEVEKQSTVAQRLHDLLELSAKNTDDVKVLKQALDELAPLRVKGGKTTVDWK  
 QKLESIESLLRATRTAGEISSELERIGTQAVGTITVRDLGTLSDQCREAANFLRQASL  
 PEGFSDIGTKLSELQAYIKYKKQFLEHFETTQPNVFQRFPLSQNITENVPARPAMDSV  
 ARLTNHLHVRGSAPHFTTWIETLPTVDPEKPTHVPAHGGAPLHRQITYSNVLEALFSL  
 CSTTLTPVPTAPGLEIATRARRGAEAAATWMDRQWPDIAQTLQDVLDTYEHTTAHANRD  
 AAFNTFLAMCVFTQIIRGASRAVTLPKLPSTAVDFPEEIVLTPRECTTLVTAMWPTLA  
 AAILRLKSYSEALGLMSRFLPLMFQALPHLTLEAQVKNGPHNTPPQLRCFAKTEAIPY  
 FPAQWQSANLEQSLWGQTDFLQICDNNQRKARVAAVTWALTIDGVVLDQLWSTFKPM  
 TAASDDTYVDLVETLHLTTFGPRGPTPRRETTTEHPPYEYQGPTGYCISGQSTTPVQA  
 SNTPVSAFEAVLGAMVFHVPIRIFLAATPKRLGQARGGMGLLTPILECVPDVEPFKSL  
 YNAPRKVPVPIETLPASLHPHDERQVFLRQAQWLSYRFTPHEAARSSTPPLLVIDPEN  
 LVTATYSSGGPANFESRPFYVMPGPYPDPWPKTLSVTSNTSVTHLSHDEICNLFTTLS  
 REHGTVOGRDIFAAAPTNTVTEQQTANPPAWETDNRLITQTETAKKPHIIPASPKARTD  
 PPVETTTTHSQGQASQHANSNVNQPQGQITSHASRNTPTAPQASSSPEKFNTQTPVRL  
 ISQSETAHINQPASGQVTEPKGIFGTYKPRVLTEPAKPANAGVASRQPEATTTVPKL  
 PINPPTARVFIGTASKLSPAVEESHGATPDHQS KIDREKYAESRPRTPHLEEGPRE  
 PHVNTPTS AHINVPSSQGQKT VHGRENPGLQTATPSAQPTASNPRIQYTLPRTDGRL  
 LHDESEVESTPTEEVKRSPKTQDVSHGPEPDDSRWTAPLGPTIEIHRLEHPQILKNIT  
 SLTVPTPRVTPIPTNIWIPLSHVNIQHEEITRAKNVLMRFIQNVRRKLQASSDALSE  
 AIARIKFLYL"

SEQ ID NO 144

CDS

complement (107637..108146)

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/note="capsid protein; ORF 65; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 65"  
 SEQ ID NO 145 /translation="MSSLRVKEPIVQGRLEHDYPNHPLVAEMNNLPQGDMSPAQYAIA  
 KRNLYVFLTAKHHYDMYMQKNGILRKDHLRGLRGKKDASSSISGVLSGSGSAAPSV  
 PVASTLGSNSFTTISGPHSLIGSMGPAPGGGGPGSVASSGIGSTSLSPSDATTLDTR  
 RSSQNKKSK"

SEQ ID NO 146 CDS complement(108152..109498)  
 /note="unknown; ORF 66; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 66"  
 SEQ ID NO 147 /translation="MASGRLPNLAEDEAACHGRGSYPahrWLDGSRLGLDLAASIRSI  
 GLCPECYVCFVTYGLGAWDGRPPKWACTLISAPSFQALNEIATGWRPDNPPKNGDVR  
 SRLHDIGRSLLEAYAWVLRCICTGVGCPSEGLSLTAVPRSAWSRYLVVSFQACCLV  
 CKTLNCRQRFPLVTCLPQHALLDLPVLRKKWNGGGCVSMQLNVPSISRRLGANLNEVSP  
 GPSDAGLLASLRELAPTVPCGNPFNALLRSLTFRALLSMSRVVLPIGESTETEISRD  
 GQKVLAYNVLFPCISLPVWSQVVARSVLEKTVAPRVVVCLECGYCLNFGRGKFETVN  
 FPPTNVFFSRDQKEKQLSICATTGRVYCSYCGGSHMRVISLFEITCVGDPYLRCVLN  
 NAAHAIRDANSLVSVVVPCLASPCATGLLKHRLVAELFYLTSSISSLSCGKCNR"

SEQ ID NO 148 CDS complement(109524..110198)  
 /note="tegument protein; ORF 67; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 67"  
 SEQ ID NO 149 /translation="MSSGKRLVDELCDLVVSYLGPSGISLDLERCQDGAPVYAKGGAV  
 PVCTVRLQHGCVYHLEFVYKFWLHKLERLAYPFAPCFVIINNGLATTLKCFCKPRDA  
 DAQFGKNLPINSDVYLERNSSVSLGQDDFMKFARLVFSGDLNVYSSMVICRTYFTEH  
 RQVLQFLVVTPKSAKRLKTLRTVFALTGHSDGLGALRRTGSVARPSGSELKDIGRGE  
 RAAMTN"

SEQ ID NO 150 CDS 110609..111982  
 /note="glycoprotein; ORF 68; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 68"  
 SEQ ID NO 151 /translation="MFVPWQLETLMRHWPSLRGLVEQSFLPGTPDGAFNSPVLHTQD  
 SLQPASSCRVCSLLFTLVRTFPPPDSEFFEDYGWLCLTCLYAPRSWTATLMVAADLLEL  
 THVYFPQCVKDGPPVYTAQSILGIDVQLHFFATRCFRPIDREQILHTSHLNFLOTET  
 GMLEGTIPGSFCFKTSWPRTKDDQQPTVACCSVGRGSHNTNRDNLDPEDLEEAFNSTN  
 AEEKPSLLGVFSATWAESQLLGSDDTQQADTHLQPSAFPTPEDADQSQGPCLMHPTLNL  
 KTKNHTASICVLCECLAAHPDAGPVLKDLRRDILENMENNVLVNRISYILNDPDSLS  
 HVRDEHLRGLIKRCSAQEIHKHFCDPVCVLNTYSHCPAVLFKCPPPEKYKCLKARLA  
 TGEFLDCNRIFDCELTQTLAVLFKGSQAKIGKTSLEIIRELGFQLRRHNIQITHPF  
 QTSNLYI"

SEQ ID NO 152 CDS 112004..112897  
 /note="unknown; ORF 69; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 69"  
 SEQ ID NO 153 /translation="MPKQPRSLASRAPYAPSVRRPDGPQSTRPASRHGSCCKSEIMQW  
 KKLVSDDTQFFSALTRRHLEGVDFLREMGTPICTSKSVMLPLNLKTIAPGRCVSLSSFG  
 HSSNMGFNCSSCTPTDRSAVSLDANALGEDSARKNSELCSVALTFYHHAQKVVQHKGF  
 YLSLLSHSMEVVRKSFTQPGLLYAHVLVLTFGHDPLPIFTVDADERLALWAVFHTRD  
 HLGETSLRLIMDNLPNYDITVDCIKQTYIMKFTPSRPNATVTVPVNSICEAVATLDC  
 TDEFREEIQRTAIINSQGLL"

SEQ ID NO 154 CDS complement(119211..119735)  
 /note="FLIP; ORF 71; similar to Kaposi's  
 sarcoma-associated herpesvirus ORF 71"  
 SEQ ID NO 155 /translation="MFPHKRLVDFGRHLEADDREAVLWLFDRPASDDTPEGFANGLCP  
 STGEPGIPLPVILLEAVFLVGRDLVSTFFLLDVGFIIERLRSSPSYFSYPKHLMLSID  
 RQLSERDVKNLVFLTGDQLGRRRNQSPFFRWLSQMEKAALVSPSNYMLVSDLLQAVS



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RRDVAKVVAANAPG"

SEQ ID NO 156 CDS complement(119794..120558)  
/note="cyclin D homolog; ORF 72; similar to Kaposi's sarcoma-associated herpesvirus ORF 72"

SEQ ID NO 157 /translation="MASVGPVPTGTIDPVLYQDRALSNLLAHEASFVTSTACYGTIQT  
EVTVMRVLGTWMRSVARAHQADASVFPLAVSILDYLECRSIPRRRRFQRLGAACLF  
LAGKIRDNLNPFKAFLCFAAEDFSVADLLKQEKSIKALRWKLEAVLPTDAIGPTLF  
KSGFTKEQLFALHSQVSVHKAIVNPATGGLPPSLVAAACALFSLGAAAPPPARLAE  
AVGVSAATLAAAESVATTREFDEDHILSNARGSS"

SEQ ID NO 158 CDS complement(120866..122212)  
/note="latent nuclear antigen; ORF 73; similar to Kaposi's sarcoma-associated herpesvirus ORF 73"

SEQ ID NO 159 /translation="MWGSRQHRSGIVSGHLRSSCRGHCGRGGTREQAGRRGRGRGT  
AAPAAAPAPPAPTTSQPVRAVAEQGHGSDTETATESRHGSSQSGSPSGSSESVIVLG  
SPTPSPSGSAPVLASGLSPQNTSGSSPASHPSPSPSPSHPGPHSPAPPSSHNPS  
NQQPSSFLQPSHDSPEPPEPPTSLPPDPSGPPQSPTPTSSPPQSPDPSGPPQSP  
TPQQAPSPNTQQAVSHTDHTGSPRPGPPFGHTSHSYTVGGWGPTRAGGVPCRLR  
CTSHNSHEDEAPERQQEQEGERQQQPARPPRPPRPPRYPIPIPYPSSEEEVPRKYRP  
QRRFYRQVLGPRIDPPRPGPWCHGVIFCNSDPYSLYRLARCLQFPGIRASSVRVLPDA  
PGSPVIPAFICITVFCQSRGTAKAVKKARRRWERHPSAPHFQASIVRMDRGLPIQH"

SEQ ID NO 160 CDS 122866..123627  
/note="R15; similar to Kaposi's sarcoma-associated herpesvirus K14 and ox-2"

SEQ ID NO 161 /translation="MSGGITLTLATLATVRCALQTHYAAVPVHSTASLGCVLTTTPH  
DVLIVTWQKQESPPVNVATYSSEAGTVVQPPFAGRVDIPEHKLTRTTLKFFNATLED  
EGCYLCIFNAFGVGLSGTACLTVYVPLSMSVTIFYPPINPTQLVCRAEASPAPSVNWT  
GVPELCEPEVFPRPNGTTLVVGRCNVTSVDPEDLENATCLVTHIGGLAAARPLDPV  
FSDPLEGTSHYVVGVVAAAALVGLTGVFLYRSM"

SEQ ID NO 162 CDS 123924..124952  
/note="G protein coupled receptor; ORF 74; similar to Kaposi's sarcoma-associated herpesvirus ORF 74"

SEQ ID NO 163 /translation="MDALNNLNLMDFLSNYSNSYSSYDDNMSYTLDTSTLCRLTV  
VFPPTVYAIICFFIFCITLFGNALVLYIFFKFKALANSVDVLMAGLCCNSLFLCASFL  
FSWLLYVAPQMLTSATCKVEIFFYLYTYFGVYIVVCISLIRCLLVFSRRPVWKHGA  
SGFLCVCVSLIVALALSANASLYRTALRHPETSEWICYEDAGEDTVNWKLRIRTTSAI  
CGFLVPFGLMVLFYGLTWCVMVKSTKLARKGAVRGVIVTVVLFILFCLPYHLNFFDT  
LLRTGFLAETCYLRDVISVAMHICSLQLQSMYSAFVPVVYSGLGLFRRRVDRDTSVFR  
CFSTSGSL"

SEQ ID NO 164 CDS complement(125057..128953)  
/note="tegument protein; FGARAT; ORF 75; similar to Kaposi's sarcoma-associated herpesvirus ORF 75"

SEQ ID NO 165 /translation="MAQRTNPRWAAAALSPREEAFIHNSDAESVLALVPEQCFSEFL  
LWLVTSPSDNFDNDDDDPALGVIWHLLAPLVNYAPLETRSAHLQGHHTISLPYGPDLN  
RQPTTRSSEIVQCLRDSDGLDRTLRLLEVGRHLSCQTRRFVADRVPPTLAALTGLTLVE  
YDVRVQRQLPVTLOSTAWRPLPERDPICAAVMLPLQRNILPLAVQASNGNSYTVSRYA  
VMARRSYSCVFQRLPCENVTHIADSFTHLHSAIQTGAGALQNILFHATLLPGGEIRSA  
LCGFYATTPSVGAFSRARHRAINTTATLHCQQLARTGTPTVLGGFLKTVHSATTSEANV  
ITTTSLSCVPQAYTFLRRSLFSQPIICLGSFEPVDGDNQSRSLYLGSAGITRITQT  
LSLAYEILEGPLFTSINRAHEPASVIGHLGALVSRGGLRFLVSQLPPTILSQLTATPD  
ISRETVDILVNKFLNVSAVVFAVLPRDTEPEPGPLDAIRRAARICGCPFAVVGETC  
EELGIQFVNDLELWNPAGAWPIROPTSAEVIATFGFDEQPVSSNWLVRPEEPEDGGEQA  
PSPTDWGLFRLASVVDQLLRCPVTGSKFVTRHVDRCNGLVAQQCEVGPLGRPLSDY

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HIVNHTSVFTDRMARVPIYRQPITRQDATERLVSPETWVTQGRGRNRWVGQCVAYGE  
 QAYKMGINA AVGARYAICEAVTNIMLAHVRRSLDITLTASVGWNPEDDQAWLLQHTLF  
 ACKELCRDLSINFAITSAGSTPCLSEELISATQQHQTVAPVPFNAVITATAEVKSSR  
 QRVTPDLKATGNLIVLVSPVPHLTQGSTFEHLCLLPSTLPDVQATHLANLFMLTEA  
 LLSRGLVVS GHVDSDGGMVVTAIEMALAGNRGLQIRIPSEETPLQWLVS ETPGVIFEI  
 QPQHVDEV RQACQNFDCRATVCGTVGQEGLSERIVISHNNDDEVYSQTLTSVAANWTSF  
 SDEQWYSWG PSFTPAQELYRKDYGCNQHN LGHLAEVCRNSELTLFATPSRPPAVAALV  
 TPGAPLPRALMAAFTNVGFDVA AVSTDDL RGGNLRGFSGLTIGGT VGIEDSYVGARC  
 AIMGLLNDPGCYGGLMAFFRRADTFSLCCGEFGFQLLGALGLLRETPHDT PGPKTPDQ  
 WDIHLEENASGNHECLWNLNLIHPQTTISIMFRVLRGLVLPGWANGRYLGVRYP RAME  
 YHLNQQRIALNYHTGNADPRMFAQHYPRNPSANSAAVAITSPDGRHLASLVDPAVTF  
 HPWQWAYVPPELADMTVSPWALAFQSLFLWCIRNRQ"

SEQ ID NO 166

PCR primer

CCTATGGGCTCCATGAGC

SEQ ID NO 167

PCR primer

ATCGTCAATCAGGCTGCG

SEQ ID NO 168

PCR primer

ATATTAAACACTCGCCGC

SEQ ID NO 169

PCR primer

ATGAGGGGCCTTTTCGTGTGC

SEQ ID NO 170

PCR primer

CTGAATCCCGCTGCCAAGGCC

SEQ ID NO 171

PCR primer

ATGTTCCCTGTCTGGTTCGTC

SEQ ID NO 172

PCR primer

TTACATCATAGCTATTGCGCG

SEQ ID NO 173

&lt;213&gt; Macaca mulatta rhadinovirus 17577

nucleotides complement(23398..23668)

SEQ ID NO 174

&lt;213&gt; Macaca mulatta rhadinovirus 17577

nucleotides 25065..25368

SEQ ID NO 175

&lt;213&gt; Macaca mulatta rhadinovirus 17577

nucleotides 25518..26525

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SEQ ID NO 176  
<213> Macaca mulatta rhadinovirus 17577

nucleotides 114979..115383

SEQ ID NO 177  
<213> Macaca mulatta rhadinovirus 17577  
nucleotides 115385..116413

SEQ ID NO 178  
<213> Macaca mulatta rhadinovirus 17577

nucleotides 131731..131926

SEQ ID NO 179  
<213> Macaca mulatta rhadinovirus 17577  
nucleotides 132333..133719  
partial terminal repeat